

REFERENCES

- <111> Franklin, Tony N.
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 Smith, John M.
 Misher, Linda E.
 Miller, David L.
 Petter, Mark W.
 Wang, Aijun
 Skeiky, Yasir A.W.
 Harlocken, Susan L.

<12> DIMENSIONS AND METRICS FOR THE
 THERAPY AND DIAGNOSIS OF BREAST CANCER

<130> 210121.419010

<140> US

<141> 2000-10-26

<160> 326

<170> FastSEQ for Windows Version 3.0

<210> 1

<211> 363

<212> DNA

<213> Homo sapien

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tgtctaaagg cgattgaagt cgtccagggg catgatgagt caccaggagt gtttttagag	180
cacctccagg aggtttatcg gatttacacc ccttttgacc tggcagcccc cgaaaatagc	240
catgctotta atttggcatt tgtggctcag gcagccccag atagtaaaag gaaactccaa	300
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ttt	363

<210> 2

<211> 121

<212> PRT

<213> Homo sapien

<400> 2

Leu Gln Thr Gln Leu Gly Pro Asn Trp Asp Pro Asn Phe Ser Ser Gly	15
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Gly Ala Ala Gln Lys Pro Ile Asn Leu Ser Lys Ala Ile His Val Val	45

Ala Arg His Arg His Ser Leu His Val Phe Leu His His Leu His Ser
 1 35 40
 Ala Tyr Arg Ile Lys Thr Phe His Asp Leu Ala Ala Phe His Asn Ser
 45 50 55 60
 His Ala Leu Asn Leu Ala Phe Val Ala His Ala Ala Phe Asp Ser Lys
 65 70 75 80 85
 Arg Lys Leu His Lys Leu His Gly Phe Tyr Trp Asn His Tyr His Ser
 90 95 100 105 110 115
 Ala Phe Ala Asp Ser Leu Lys Gly Phe
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 aaaaatctnc tgccttttc aaggaacctat cccatccatt cctnaacaaa aggcctgcen 840
 ttcttccccc agttaactnt ttttnttaa aattcccaa aaangaacen cctgctggaa 900
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[illegible]

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aggatcaact	gctctaatca	ccatgacatg	aggtaaccac	caaaccatca	agcgctaaac	240
agacagaatg	tttccactcc	tgatccactg	tgtgggaaga	agcaccgaac	ttaccacact	300
gggggcoctg	ntcanaanaa	aagcccactg	ccccgggtnt	noctttnaac	cggaacgaat	360
naacccacca	ccccacanc	tcctctgttc	ntgggcoctg	catcttgttg	octentntnc	420
tttnggggan	acttggggaa	ggtaccccat	ttctttgacc	cnctnanaaa	acccctngtg	480
ccctttgccc	tgattcnctt	gggccttttc	tcctttccct	tttggggtgt	ttaaattccc	540
aatgtccccc	gaacccctct	ctncttgccc	aaaaactacc	taaatttctc	ncctangntt	600
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ntatnntggn	ccctnaaaaa	nnctctnccc	caacttgccc	gaatttggtt	ggttttttcc	720
notgggggaa	accttttcaa	tttcccccct	ggccggcccc	ccctttttcc	cccttttnga	780
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aaantccctt	ccccnaaaaa	ggttgcttag	ccccngtccc	ccactccccc	nggaaaaaat	960
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- *210* 6
- *211* 950
- *212* DNA
- *213* Homo sapien

$\frac{d}{dt} \left(\frac{\partial L}{\partial \dot{x}} \right) = \frac{\partial L}{\partial x}$

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<401> 5

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aaactatctg aactatctga aactatctgt agtccatctt ccttttcttt cccctctctc      127
ttttttttta aatcaactta aactatctgt taactcaaat tcttcttttt cgtgttatga      177
ttttttttta ttgttggaat ttcttggaat ttcttttaatt ttcttttttt atgtgttttt      227
aggaatctgt gatgttgacg ttcttcagag cagttatctc ttcttcagag atgtgttttt      277
tttttttttt cccctgttta ccaaggagat ttatctatct ttgttttttt ccttttcttt      327
aaaaacttta nctncttaagg ctgaattgtt cctctcactt cagttatctg gatggggtaa      377
ttttttttta tttaaaactt taagagaggt aggttttttt ttgttttttt ctactgtgag      427
cactctacaa ggagctgggg ctactcactt ggaggttggg ttgttttttt ttgttaangga      477
catcaaaagg aaaaacnngg ttgttgcctg tttaaaacaa aactgtatct ccagctctaa      527
gatctgtgtt tgaatttctc tctnctctct taaaattgat gctcaacttc tcttttctca      577
accagatctg cctgacaact cccatactta aaaaaaaan aanaattggc ccgaacccna      627
accaataaaa atgggggagg ttggttganc nncctgaac aaaaataatg gatcccccgg      677
gtgtcaggaa ttcaattcct ccttatctat acccccaan nggggggggg ggccgtctnc      727
cattncctct ttatttattt tttnncctcc cccccggant ccttttttaa ctcgtgaaag      777
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<111> 9
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 <113> DNA
 <114> Homo sapien

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agaactccat agtgaggtaa aagcctgggg ctttttcagag aaggggaggt tatgggtttt      180
ccaattatat aagtcagaag tagaaagaag ggacataaac caggaaaggg gtggacactt      240
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acggtttgaa ccaagagaca atgggtgaty agcctacag ggacataaac gaggagacat      360
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aagaggttct gttcaggaa gctttggaag taaaattatg aatggtgcat gaatggagct      540
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[illegible]

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[illegible]

100

[illegible][illegible]

- *221* misc. feature
- *222* [1]... [571]
- *223* R = A, T, C or G

- 100 - 14

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aaatacatat	ttcataattt	aggttagagg	gattttacag	gttcaatttt	aaattcttta	180
aaacagtttt	taagtatatt	ggaaacagat	atattctttt	tcttgggcag	tattaaactt	240
atagcaaat	tgtgtctggg	ggactgctgg	taactgtctt	taacagttgc	aaatcaaggc	300
atttgcaac	aaataaaaa	aatttttttg	tttaatttga	aaatggagtg	gataaaaagt	360
gtttggtag	gtgtgtgtat	atagttttta	atggtttatt	gaactctttt	aaattgcaat	420
tatgtggggg	ggggtttttg	atagaaaagt	ttttantcac	aaagtacacg	ggactttttt	480
cttttggrna	ctgagctaaa	aagggtgnt	tttgggttgg	gggagatga	aggtcacag	540
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46-25

[illegible]

1. The first step is to identify the key components of the system. This includes understanding the hardware, software, and data involved.

2. The second step is to define the requirements. This involves determining what the system is intended to do and what it must be able to do.

3. The third step is to design the system. This includes creating a detailed plan of how the system will be built and how it will be tested.

4. The fourth step is to implement the system. This involves building the system according to the design and testing it to ensure it meets the requirements.

5. The fifth step is to maintain the system. This involves keeping the system up-to-date and ensuring it continues to meet the requirements.

1. *Pharmaceutical industry* – The pharmaceutical industry is the largest of the three industries, with sales of \$10.5 billion in 1997. It is the only industry that has not experienced a decline in sales since 1990. The industry is dominated by a few large firms, with the top five firms accounting for 40% of sales. The industry is also characterized by high R&D expenditures, with the top five firms accounting for 60% of total R&D.

Trial	Control	MCI	AD
1	95	85	75
2	95	85	75
3	95	80	70
4	95	78	68
5	95	75	65

[illegible]

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<213> Homo sapiens
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<400> 17

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tgccttagag	gggggaagt	caatggggt	ctcaccctat	cctattgca	tgggtggtag	180
gattgggggt	tgggggggt	ttatgaccc	ggtctctctg	cgggttaaca	cctgggtgt	240
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<210> 18
<211> 262
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<213> Homo sapien
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*223*  +  1 = A, T, C or G

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[illegible]

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<411> 14

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aaatctgaga	taaaatctctc	taaaatctctc	caaaacacac	aatctctctc	caaaacacac	181
atttatgata	tataaaacac	ggttcgagct	ggttcgagct	ggttcgagct	taaaacacac	241
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tgggactgag	taaaatctct	ggttcgagct	taaaatctct	taaaatctct	ggacgtcggt	180
tgaaaatct	caaaatctct	acgatcgagc	tcttcgagct	ggacgtcggt	ggacgtcggt	240
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<211> 21

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<213> Homo sapien

<220>

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<222> (1)...(208)

<223> n = A,T,C or G

<411> 21

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gttctctgag	aaaaggggac	gagctctctc	gactattggt	attctctctc	ggtctctctc	180
caactagtag	ttttgagagc	attctctctc	attctctctc	attctctctc	attctctctc	240

<211> 22

<212> DNA

<213> Homo sapien

<221>

<221> misc_feature

1. $\frac{1}{2}$ 2. $\frac{1}{2}$ 3. $\frac{1}{2}$ 4. $\frac{1}{2}$ 5. $\frac{1}{2}$ 6. $\frac{1}{2}$ 7. $\frac{1}{2}$ 8. $\frac{1}{2}$ 9. $\frac{1}{2}$ 10. $\frac{1}{2}$

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540
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620

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60
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[illegible][illegible][illegible][illegible]

<210> 33
 <211> 477
 <212> DNA
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450 328

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ataacttttt	caacgtaagg	tctcttgctt	gttagtgtag	tgtgggttata	ttaaaatttt	180
tagttattat	tttttatcca	cttttccact	agaaagtcac	tattgattta	gcacacatgt	240
tgatctcatt	tcattttttt	tttttatagg	caaaatttga	tgtctatgcac	caaaaaaac	300
caagcccatt	attttttttt	ccccggaaat	ctgaaaattg	caggggacag	aggggaagtt	360
tcccattaaa	aaattgtaaa	tatgttcagt	ttatgtttta	aaatgcacaa	aacataagaa	420
aatttgtgtt	acttgagctg	ctgattgtaa	gcagttttat	ctcaggggga	actacta	477

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<210> 34
<211> 631
<212> DNA
<213> Homo sapien
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4 3

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atatatatat	attcaaaaga	agtaaaaatgt	atatgatcat	ataatttota	gttatagaa	180
tgaattaaag	tttatggcat	tgggtaggg	catattttaa	ttagagjafa	aaagaaaaat	240
tgggtatttt	tagaatgura	tacatgtttt	tttaaaaaat	tcaatatatta	ttttagatatt	300
tatataagaa	tgggaatttt	attaaaaattt	actaaaaaa	tattttttga	ggagagaaaa	360
attgataat	ttgcatatt	tacagtgata	caagtgatg	tcaagttaaa	taattatatga	420
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ggtttatcat	tttggaatt	tttagatata	attatttttt	tttggttttt	gattatgttt	540
aaattatttt	tttttttttt	atttttgggt	tacatttttt	tttttttttt	atttttttttt	600


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cttctctctt cttctctctt cttctctctt cttctctctt cttctctctt  1-
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cttctctctt cttctctctt cttctctctt cttctctctt cttctctctt  120
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<211> 39
<212> DNA
<213> Homo sapien

```

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<220>
<221> misc_feature
<222> (1)...(39)
<223> n = A,T,C or G

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<411> 39
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cttctctctt cttctctctt cttctctctt cttctctctt cttctctctt  153
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<210> 39
<211> 585
<212> DNA
<213> Homo sapien

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<220>
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<222> (1)...(585)
<223> n = A,T,C or G

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ttagaaatgc atatgcctct ataatacaca actgatttag aagctattac aattaaaaag  180
ttggcccccg cgtggtgggg ggtgggtgag gctgtgaatc ctggcacttt ggtaggccga  240
ggtacgggga ttaggggggt gggagttttaa gacatcctg gctaacaggg tgaaggttca  300
tctctactaa aaatacgaaa aaattacccc gggtggtggg cgggggcctg tagtcccagc  360
tactccggag gctgaggtag gagaatgggg tgaacccagg acacggaggt tgtagtgtgc  420
gaaatcaggg ttagtgcctt ccaactgggg ggaacgggaa agantccgg tautcaaaaa  480
agaaaaatat tctatatttt tctacttata tcttaantta cagacaaatn cctcttgcta  540
cctctttagc atttatttca cctacttact atagggaan nttaa  585

```

```

<210> 41
<211> 477

```

Figure 1 shows 16 small diagrams arranged in a 4x4 grid, labeled 1 through 16. Each diagram illustrates a different configuration of a 2x2 grid of squares. Some squares are shaded, some are outlined, and some are empty, representing various states or transformations of the grid.


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<210> 44
<211> 144
<212> DNA
<213> Homo sapiens

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cagatggac	cttcaggaga	ttgtgtacac	tagaaaagaa	ctgttgagt	ttcttaattt	180
atataagcag	aaatctggag	aagagtcata	ggaatggata	ttaaaggt	gttagataatgg	240
gggaagggaat	atagagtttg	atcaggtggg	acttattgat	ttgaacccac	taagtacaga	300
ttctgttttt	gattgttgag	ctcaggtgag	taaaaaaggt	tttaattggt	ctaatagttt	360
attgtgttgg	ctagctgaaa	tatggataaa	agatggccca	ctgtgagcaa	gttgaaatg	420
ccgtgtctct	ctcaatttaa	tgtaacggaa	gggatacaaa	agtttaggga	ganttggatg	480
ctgagkttag	atgggttaat	ttgtagcata	ccwtctcag	ctgggaggt	ccagagcata	540
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<210> 44
<211> 144
<212> DNA
<213> Homo sapiens

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<220>
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<223> n = A,T,C or G

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<401> 44

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cagatggac	cttcaggaga	ttgtgtacac	tagaaaagaa	ctgttgagt	ttcttaattt	180
atataagcag	aaatctggag	aagagtcata	ggaatggata	ttaaaggt	gttagataatgg	240
gggaagggaat	atagagtttg	atcaggtggg	acttattgat	ttgaacccac	taagtacaga	300
ttctgttttt	gattgttgag	ctcaggtgag	taaaaaaggt	tttaattggt	ctaatagttt	360
attgtgttgg	ctagctgaaa	tatggataaa	agatggccca	ctgtgagcaa	gttgaaatg	420
ccgtgtctct	ctcaatttaa	tgtaacggaa	gggatacaaa	agtttaggga	ganttggatg	480
ctgagkttag	atgggttaat	ttgtagcata	ccwtctcag	ctgggaggt	ccagagcata	540
caactttgat	caaggttttg	cgaaatggat	ttgtgatggc	ggcaactact	aa	592

```

<210> 45
<211> 567
<212> DNA
<213> Homo sapiens

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<220>
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<223> n = A,T,C or G

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ggctggttgg	ctttgaaag	atggaaatcc	tttaggacat	gttagaaaag	ctttcttggc	180
gaacagtttg	ttctgggag	aaagctcacc	aagatgcccc	ttgaaaggg	taycgtgtat	240
ttgggagaga	ctgataaggt	gtcttctgat	gattgttttg	cttggacagt	gacaaaagat	300
atgcaaaaga	agtgcgaact	agaagtcaag	cttctgtagc	aaattattgt	agactcttat	360
ttatacttga	agaaatgata	gcaaaaggtg	ggtaatttaa	gaataaggtg	gtttgtaatt	420
gagaggtatg	ttctaggga	aaagacatga	tggatagttt	tataatggga	actaatcaag	480
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ctatacttga	gtttctata	atgttccc				567

```

<210> 46
<211> 567

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1. *Chlorophyll *a** and *Chlorophyll *b** were determined by the method of Arar and Collins (1971).

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<210> 47
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<212> DNA
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<220>
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<223> T = A,T,C or G
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<400> 47

- 210. 45
- 211. 591
- 212. DNA
- 213. Home Station

[illegible][illegible]

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tttttttt tttttttt tttttttt tttttttt tttttttt tttttttt 24
tttttttt tttttttt tttttttt tttttttt tttttttt tttttttt 30
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<210> 40

<211> 454

<212> RNA

<213> Homo sapien

<400> 41

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aattgggtta tttttttt tttttttt tttttttt tttttttt tttttttt 180
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aattgggtta tttttttt tttttttt tttttttt tttttttt tttttttt 300
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aattgggtta tttttttt tttttttt tttttttt tttttttt tttttttt 420
tttttttt tttttttt tttttttt tttttttt tttttttt tttttttt 480

```

<210> 50

<211> 463

<212> RNA

<213> Homo sapien

<400> 50

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aattgggtta tttttttt tttttttt tttttttt tttttttt tttttttt 180
tttttttt tttttttt tttttttt tttttttt tttttttt tttttttt 240
aattgggtta tttttttt tttttttt tttttttt tttttttt tttttttt 300
tttttttt tttttttt tttttttt tttttttt tttttttt tttttttt 360
aattgggtta tttttttt tttttttt tttttttt tttttttt tttttttt 420
tttttttt tttttttt tttttttt tttttttt tttttttt tttttttt 480

```

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<211> 399

<212> RNA

<213> Homo sapien

<400> 51

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aattgggtta tttttttt tttttttt tttttttt tttttttt tttttttt 180
tttttttt tttttttt tttttttt tttttttt tttttttt tttttttt 240
aattgggtta tttttttt tttttttt tttttttt tttttttt tttttttt 300
tttttttt tttttttt tttttttt tttttttt tttttttt tttttttt 360
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 ctgattcagc atcnnagaa acntcttgact ctgattctag aggaactgagt gaattgaa 180
 ttgaggtggtat gctcctctcag gctacatcat tatagaagca aagtagaana atcnnngtttg 240
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 <213> Homo sapien.

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 <223> n = A,T,C or G

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<210> 14
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 <213> Homo sapien.

<220>
 <221> misc_feature
 <222> (1)...(112)
 <223> n = A,T,C or G

<400> 84
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<210> 85
 <211> 225
 <212> RNA
 <213> Homo sapien.

<400> 86
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 ctgattcagc atcnnagaa acntcttgact ctgattctag aggaactgagt gaattgaa 180
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1. *What is the main purpose of the study?*
 2. *What are the research objectives?*
 3. *What is the research methodology?*
 4. *What are the findings of the study?*
 5. *What are the conclusions and recommendations?*

• 400 •

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120
150
211

<400> 59

60
120
180
208

450, 60

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 <212> DNA
 <213> Homo sapiens

<400> 62

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 <212> DNA
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<400> 63

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<400> 64

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- >412. RNA
- >413. Home sapien

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300

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180
240
244

- * 210 - 77
- * 211 - 254
- * 212 - DNA
- * 213 - Home center

Figure 1 shows a series of nine line drawings arranged in a 3x3 grid, illustrating the development of a child's drawing of a person from age 3 to age 7. The drawings are labeled as follows:

- Top row: 3 years, 3 years, 4 years
- Middle row: 4 years, 4 years, 5 years
- Bottom row: 5 years, 6 years, 7 years

The drawings show a progression from simple stick figures to more detailed representations with features like hair, clothing, and facial features.

the 1990s, the number of people in the world who are illiterate has increased from 1.2 billion to 1.5 billion. The number of illiterate people in the world is expected to increase to 1.7 billion by the year 2015. The number of illiterate people in the world is expected to increase to 1.7 billion by the year 2015. The number of illiterate people in the world is expected to increase to 1.7 billion by the year 2015.

1. *Staphylococcus aureus* (ATCC 12228) was grown in tryptic soy broth (TSB) (Difco, Franklin Lakes, NJ, USA) at 37 °C for 24 h. The bacterial suspension was centrifuged at 1000 g for 10 min and washed with distilled water. The cells were resuspended in distilled water and adjusted to a concentration of 1×10^8 CFU/ml.

100

0	1	2	3	4	5	6	7	8	9	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T	U	V	W	X	Y	Z	[\] ^ _ ` { } ~								
a	b	c	d	e	f	g	h	i	j	k	l	m	n	o	p	q	r	s	t	u	v	w	x	y	z	{	}	~		!	"	#	\$	%	&	'	()	*	+ , - . / : ;						
:	@	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T	U	V	W	X	Y	Z	[\] ^ _`	{		}	~		!	"	#	\$	%	&	'	()	*	+ , - . / : ;
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:	@	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T	U	V	W	X	Y	Z	[\] ^ _`	{		}	~		!	"	#	\$	%	&	'	()	*	+ , - . / : ;
:	@	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T	U	V	W	X	Y	Z	[\] ^ _`	{		}	~		!	"	#	\$	%	&	'	()	*	+ , - . / : ;
:	@	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T	U	V	W	X	Y	Z	[\] ^ _`	{		}	~		!	"	#	\$	%	&	'	()	*	+ , - . / : ;
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:	@	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T	U	V	W	X	Y	Z	[\] ^ _`	{		}	~		!	"	#	\$	%	&	'	()	*	+ , - . / : ;
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:	@	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T	U	V	W	X	Y	Z	[\] ^ _`	{		}	~		!	"	#	\$	%	&	'	()	*	+ , - . / : ;
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:	@	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O																														

- * Z110 - 706
- * Z110 - 406
- * Z112 - RNA
- * Z114 - Home Science

Figure 1. Schematic representation of the experimental design. The subjects were divided into two groups: the control group and the experimental group. The control group was divided into two subgroups: the control group and the experimental group. The experimental group was divided into two subgroups: the control group and the experimental group.

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ataattatta	gtagtacgg	taggaga				327

- 210. FI
- 211. FI
- 212. DNA
- 213. Home studies

Figure 1. Schematic diagram of the experimental setup.

[illegible]

Figure 1 illustrates the experimental setup. A subject is seated at a table, looking at a video screen. A video camera is positioned above the screen to record the subject's hand movements. A light source is positioned to the left of the screen. A target is positioned on the screen. The subject's hand is positioned near the target. The diagram shows the relative positions of the subject, camera, screen, light source, target, and hand.

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101

10

11

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[illegible][illegible]

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1. *What is the purpose of the study?*
 2. *What are the research questions?*
 3. *What is the study design?*
 4. *What is the study population?*
 5. *What are the inclusion and exclusion criteria?*
 6. *What are the variables?*
 7. *What are the data collection methods?*
 8. *What are the data analysis methods?*
 9. *What are the results?*
 10. *What are the conclusions?*
 11. *What are the limitations?*
 12. *What are the implications?*
 13. *What are the recommendations?*
 14. *What are the references?*
 15. *What are the appendices?*
 16. *What are the acknowledgments?*
 17. *What are the funding sources?*
 18. *What are the conflicts of interest?*
 19. *What are the ethical considerations?*
 20. *What are the contact details?*

[illegible]

4/20/17
 JEFFREY D. GREGG JR.

- * 210 - 108
- * 211 - 26
- * 212 - DNA
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213. Primer for amplification from breast tumor cDNA

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[illegible]

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212-131
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 Gly Ile

214-131
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 214-133
 214-134
 214-135 Artificial Sequence

215-131
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216-131
 216-132
 216-133
 216-134
 216-135 Xaa = Any Amino Acid

217-131
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 1 5 10 15
 Val Gln Gly His Asp Glu
 20

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 218-132
 218-133
 218-134
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219-131
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gacatgtgct	gtgttgactc	aaggttcaat	ggatttaggg	ctatgctttg	ctaaaaaagt	240
gcttgaagat	aatatgcttg	ttaaaagtca	tcacattctt	ctaattctca	gtaccaggg	300
acacaaatac	ctggggaagg	cggcaggga	ctctgtctag	gaaagccagg	tattgtccaa	360
gattctctcc	catgtgatag	ctgagatat	ggcttcattg	gaaggctaag	acctgaatgt	420
ccccagagac	gaattcccc	agccagacat	ccccagagac	gacacacgaa	aagggtctgt	480
gtgaggagg	attagtaaaa	gagggaagac	tctttgcagt	tgaggtaaga	ggaaggcaac	540
tgctccctgc	tgtcccttgg	ccaatagaat	gtcttgggtg	aaacccagat	tgtatgttct	600
attactgag	ataggagaac	atctctctat	ggctggaggt	gagacacgct	gagggaata	660
ccgctcttca	atggaacgag	acttttgtat	aactccacat	caaggacag	caattttctt	720
taaaatttat	tatgacacag	agacttttct	tcacatttca	ctctggaac	ctctccacac	780
attacattat	tggtctgaca	caacccctct	tcaggaatgg	ctagataaat	tatcaataaa	840
tattgagga	actaagagac	caatctctct	gtagttctca	ctctctctca	gagaggttct	900
ctgaggttca	attctctctc	tctctctctc	attctctctc	ctctctctct	ctctctctct	960

[illegible]


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TCTGATGTTT TCTGATGTTT TCTGATGTTT TCTGATGTTT TCTGATGTTT
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+210 > 144
+211 > 144
+212 > DNA
+213 > Homo sapien

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TCTGATGTTT TCTGATGTTT TCTGATGTTT TCTGATGTTT TCTGATGTTT
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+210 > 148
+211 > 111
+212 > DNA
+213 > Homo sapien

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+400 > 145
AGGCAATTTAC GAGCAATTTAC CAAAAAAGAG CAAAAAAGAG CAAAAAATATC AGCAATTTAC
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+210 > 146
+211 > 585
+212 > DNA
+213 > Homo sapien

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+400 > 146
tagcatgttg agcccagaca cttgttagaga gaggaggaca gttagaagaa gaagaaaagt 60
tttaaatatg tgaagattac tataagaaaag ctttggtttt ggatgagact tttaaatatg 120
cagaggatgc ttgcagaaaa cttcataaat atatgcaggt gattcattat ttctccttag 180
aaatttagtg atatttgaaa taatgccccaa acctaatctt ctcttgagga aaactattct 240
acattactta agtaaggaat tatgaaaagt ttcttttttag gtatagtctt tcttaattgg 300
gtttgacatt gcttcacagt gctctgtttt ttgtccataa tcgaaaagtaa agatagctgt 360
gagaaaaacta ttacctaaaat ttggtatggt gttttgagaa atgtcattat agggagttca 420
ctgtgtgtgt tttaaatatc ttgtgtatct ataattgagc taattataaa aactcttttg 480
agacatatct taaattgtct ttctctgtta tactgagat gatgtttctt cagcattttt 540
ctctcgaatt gggacatttg ctactgtgto ttggttcaca tgcta 585

```

+210 > 147
+211 > 579
+212 > DNA
+213 > Homo sapien

+210 >
+211 > misc_feature
+212 > 1..100
+213 > A,T,C,G

[illegible][illegible][illegible]

• 23. Home again.

tgagacatttg	ttaacacatct	actatcttgt	gacattctaa	taataagcaat	tcttaactagt	60
gngagatggt	aaatcaatgt	gggtttgggc	tgcattcttc	taatgataag	tgatatttaag	120
cttttttttaa	atatgcttgt	tgaccacatg	catatcatct	tttgagaagt	gtctgttcat	180
atctctttgac	caatttttaa	tttttttata	ttgtaaattt	gtttaatttc	cttaacagatg	240
ctggacacagc	tatcaa					255

• 213 • Homo sapien

ttacggctgga	aaaactgtgga	ggcgaagctg	ggatcaactt	tccattctaa	cttgagagga	60
ggaagattca	agtcaggca	aggggtgggg	ggtagaagt	ggcaacaga	aatttcagct	120
ggacactgt	caacacata	ggaggacag	aagggctgg	ctctcagg	cagatgaag	180
aaaggagaa	tgttttggt	gcaacaaag	gttagagact	cccatctttg	aaggaagggc	240
tcttttgtt	ttcttactt	cctctttctg	ctagagact	ggttatagac	aaggtctctt	300
caagatttg	caactta					360

1. *Chlorophyll a* and *Chlorophyll b* were determined by the method of Arar and Collins (1971) using a Shimadzu 1601 UV-Visible Spectrophotometer.

<410> 151
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 <412> DNA
 <413> Homo sapiens

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ctctctctc cctctgaga gctctcctc cctctctc cctctctc cctctctc	180
gctctctc cctctgaga gctctcctc cctctctc cctctctc cctctctc	240
gctctctc cctctgaga gctctcctc cctctctc cctctctc cctctctc	300
gctctctc cctctgaga gctctcctc cctctctc cctctctc cctctctc	360

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 <411> 311
 <412> DNA
 <413> Homo sapiens

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ctctctcagg ttgattctgt cctctctctt cagctctcga atgctctctt cgtctctctt	180
tggtctctct cctagtgaa cctctctctt cctctctctt cctctctctt cctctctctt	240
gaggtctctt cagagattgt cagctctcga tcagagctag cgtctctcga cgtctctctt	300
cagagctctc	311

<410> 153
 <411> 332
 <412> DNA
 <413> Homo sapiens

<410> 153	
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gttgatctct ggtctctctt ttctctctc agagtgaggaa cttctctctc catgtttgat	180
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<410> 154
 <411> 345
 <412> DNA
 <413> Homo sapiens

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 <423> n = A,T,C or G

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ctctctctc cctctgaga gctctcctc cctctctc cctctctc cctctctc	180

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Figure 6

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AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 1
AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 12
AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 24
AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 36
AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 48
A 60

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<21> 167
<211> 42
<212> DNA
<213> Homo sapien.

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<4> 167
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AGTTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 24
AGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 36
AGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 48
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AGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 72
AGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 84
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AGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 108
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AGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 132
AGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 144
AGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 156
AGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 167

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<211> 167
<211> 42
<212> DNA
<213> Homo sapien.

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<221>
<221> misc feature
<222> 1... 500
<223> n = A,T,C or G

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<400> 167
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ggagtagttc cctgctaagg gagggtagac tgttcaacct gtctctgctc cggcctccac 180
tatagcagat ggcagcagga gtaggagaga gggaggtaag agtcagaagc ttatgtttgt 240
tatgggggga caagccrtat cgggggcagc cragtattata ggggacatn tagwyartw 300
agntagcacc caaagcgggg gaggtncccc atatgggttg acctgcaggg ggcgcgatta 360
gtgattagga tgtgagcccc agacagcat agcaacaagg acctaaactc agatcctctg 420
ctgattactc caatgaatt attgtattta ttttaacaat ttgagttatg aggcattatta 480
ttaggtccat attacctgga 500

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<210> 167
<211> 355
<212> DNA
<213> Homo sapien.

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<411> 167
tccatgagac ggtgagagaa gctggaatt ccaagacttt cggaggggaa ggggagagaa 60
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tccatgagac ggtgagagaa gctggaatt ccaagacttt cggaggggaa ggggagagaa 180
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tccatgagac ggtgagagaa gctggaatt ccaagacttt cggaggggaa ggggagagaa 480
tccatgagac ggtgagagaa gctggaatt ccaagacttt cggaggggaa ggggagagaa 500

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[illegible][illegible]

the 1990s, the number of people in the world who are illiterate has increased from 1.2 billion to 1.5 billion. The number of illiterate people in the world is expected to reach 1.7 billion by the year 2015. The number of illiterate people in the world is expected to reach 1.7 billion by the year 2015. The number of illiterate people in the world is expected to reach 1.7 billion by the year 2015.

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[illegible]

- * 210 - 173
- * 211 - 701
- * 212 - 10A
- * 213 - Home Services

<400> 173

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tgotttagott	catgtaagoc	atctctgtatt	catttagaaat	aagaacaatt	ttattctgtog	240
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cattgctatt	ccattctatt	ccagagacaa	gtcaacttat	gcagagaaatc	aaactgcaag	660
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- 210 - 174
- 211 - 701
- 212 - 1102
- 213 - H. M. - 701

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+210> 178
 +211> 786
 +212> DNA
 +213> Homo sapien

+40> 178
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 +211> 786
 +212> DNA
 +213> Homo sapien

+40> 178
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 gtttttttt aggtttttt aggtttttt aggtttttt aggtttttt ttttttttt 360
 cagtctaac ttttttttt gtttttttt gtttttttt aggtttttt ttttttttt 420
 ttttttttt atatatatgt atgtatgtat gtttttttt aattttttt ttttttttt 480
 gtttttttt ttttttttt tagtttttt atttttttt ttttttttt atttttttt 540
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 gtttttttt gtttttttt ttttttttt atttttttt ttttttttt ttttttttt 660
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 atttttttt atttttttt ttttttttt gtttttttt atttttttt ttttttttt 780
 atttttttt atttttttt ttttttttt gtttttttt atttttttt ttttttttt 840

+210> 178
 +211> 786
 +212> DNA
 +213> Homo sapien

+40> 178
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 aggtttttt aggtttttt aggtttttt aggtttttt aggtttttt ttttttttt 360
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60
120
178

60
220
180
240
300
360
367

[illegible][illegible]

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*210* 193
*211* 372
*212* DNA
*213* Homo sapien

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[illegible]

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120
180
240
300
360

6

1. *Journal of the American Medical Association*, 1997; 277: 1033-1038.

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1000

[illegible]

Table 1. *Salmonella* serotypes and phage types isolated from the 1990-1991 and 1991-1992 seasons

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100

423-424

5.3.2. **RNA**

2014, 11:10

* 221, m. 15

E. coli O157:H7 was detected in 10% of the samples.

• 2017 年 11 月 25 日

[illegible]

Figure 1. Schematic representation of the experimental design. The subjects were divided into two groups: the control group (CG) and the experimental group (EG). The CG was divided into two subgroups: the control group (CG) and the control group (CG). The EG was divided into two subgroups: the experimental group (EG) and the experimental group (EG). The CG was divided into two subgroups: the control group (CG) and the control group (CG). The EG was divided into two subgroups: the experimental group (EG) and the experimental group (EG).

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130
240
300

<210> 203

<211> 241

<212> DNA

« 213 » Homo sapiens.

450. 203

60
225
230
240

1. *Chlorophyll a* (Chl *a*)

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1. *Chlorophyll a* and *Chlorophyll b* were determined by the method of Arar and Collins (1987).

The diagram illustrates the experimental design. It shows a sequence of three events: 'Stimulus' (a face), 'Response' (a button press), and 'Reward' (a coin). The sequence is labeled 'Stimulus', 'Response', and 'Reward'.

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 22-12

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[illegible]

1. *Staphylococcus aureus* (ATCC 12228) was grown in tryptic soy broth (TSB) (Difco, Franklin Lakes, NJ, USA) at 37 °C for 24 h. The cells were washed with phosphate buffered saline (PBS) (pH 7.4) and resuspended in PBS. The cell suspension was adjusted to a concentration of 1×10^8 CFU/ml.

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421. Homo sapien.

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[illegible]

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[illegible]

- **Chlorophyll** is the green pigment that captures light energy
- **Chloroplasts** are the organelles where photosynthesis takes place
- **Stroma** is the fluid-filled space surrounding the thylakoids
- **Thylakoids** are the membrane-bound structures where the light-dependent reactions occur

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140

- * 210 * 230
- * 211 * 257
- * 212 * DNA
- * 213 * Home sapien

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| atgtttgtta | atogaaggt | aaatgttga | gaggaatgat | aaatattat | aatgaatga |
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[illegible]

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 <223> n = A,T,C or G

<400> 237

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 tgcctgtctta cccatctcaa aagactgcga aaatcacaac agttgttga acagaaatat
 aagaaatatc cctcattctt ctctcttcaa tctacttcac tctcaattca ttagtaata
 atcgttttca gaaaacaaa cacttctagt tctcactcat aagcttgaat tgaacaa'aa
 gaacacaaag acacagggag ggaacatca caaacacagg ccttcacagg agtctgggag
 atgagtcagt cta

60
 120
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 300
 360
 420

<210> 238
 <211> 492
 <212> DNA
 <213> Homo sapien

<220>

<221> misc_feature
 <222> (1)... (492)
 <223> n = A,T,C or G

<400> 238

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 atactcagagt gattagaaga agtggacaga gctacccaaag ttaaacatat gacagatcaa
 aaaaatatag cacttgctaa cccacactac agggagggaaa taaggaaatc aatagcatat
 tgtgttatth tgaatgcac gaaatctctt anaanaaaaa ataac'aaag aaaaacagaa
 aatttccttca aatgttcaat cctctagaag aatttaacaa aaacatatat tcaatgaatt
 caaaaatctt agcaggttca agcacaacaa atcaccagac agactaatcc cactttagat
 tctcagatcaa actaaacag aagaaatccc aatgaacatt caattctaa gaa'aaacaa
 taattctctca

60
 120
 180
 240
 300
 360
 420
 480
 540

[illegible][illegible][illegible]

| | | | | | | |
|------------|------------|------------|------------|-------------|------------|-----|
| tgtatcgagc | tagtggctct | cccatgtgat | agtctgaaat | atagctccat | gggatgagag | 60 |
| gctgtgctcc | agcccgacac | ccgtaaaagg | tctgtgctga | ggtggattag | taaaagagga | 120 |
| aagcgtttgc | gttgagatag | aggaaggcca | ctgtctcttg | cctgccccctg | ggaactgaat | 180 |
| gtctcggtat | aaaaccogac | tgtacatttg | ttcaattctg | agataggaga | aaaaccaccc | 240 |
| tatggcggga | ggcgagacat | gttggcagca | atgctgcctt | gttatgcttt | aactcaagaa | 300 |
| tgtttggggc | gagggaaaca | taaactctgc | ctacgtgcac | atccaggcat | agtaacctcc | 360 |
| tttgaaacta | attatgacac | agattccttt | gtccacatgt | tttttgcctg | acctctctct | 420 |
| tattatcacc | ctgctctctc | accgcattcc | ttgtgctgag | ataatgaaaa | taatatcaat | 480 |
| aaaaacttga | ccgaactcgc | agaaccacta | gtcgatata | | | 519 |

- *220. $\bar{A} \vee \bar{B}$
- *221. $\bar{A} \vee \bar{B} \vee \bar{C}$
- *222. $\bar{A} \vee \bar{B} \vee \bar{C} \vee \bar{D}$
- *223. $\bar{A} \vee \bar{B} \vee \bar{C} \vee \bar{D} \vee \bar{E}$

[illegible]

<210> 246
 <211> misc_feature
 <212> DNA
 <213> Homo sapien

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atgaagtttt agaccatcat tctcagcaaa ctacaaaggg aacagaaaaa caaacattgc
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tgtaacaaa atgcattgtt tgcacatgta ccccagaact taaagtgtta ataaaaaaat
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aa

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<210> 246
 <211> 474
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...4421
 <223> n = A,T,C or G

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<401> 246
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atgtttctca tcttaagctg gattgaara atgagaacac atggacacag ggaaggaa
atcacacagt ggggcttggt gttgggtagg ggtctaggga agggatagga ctaggagaaa
tatctaattg agatgaaggg ttgatgggtt cagcaaaacca ccatgacacg tgnataccta
tgtaacaaa atgcattgtt tgcacatgta ccccagaact taaagtgtta ataaaaaaat
taagaaaaaa gttaagtatg tcatagatac ataaaaatatt gtanatatgt aagggtgccc
aa

```

<210> 247
 <211> 474
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...4474
 <223> n = A,T,C or G

```

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gtttgggaa ctgacattat ttgggtatga gtaacagaga aaaaagtgga aaaggtttta
aatatattta atattattt ttagaatttt cactcttgga gttttcaat aactatctag
ttttttttt ctatatttt cactttttt atatttttt taatttttt ttgtttttt
taatttttt ttatttttt atttttttt atttttttt ttgtttttt gattttttt

```

1. *Pharmaceutical industry* – The pharmaceutical industry is the largest of the three industries, with sales of \$10.5 billion in 1997. It is the only industry that has a significant presence in all three markets. The industry is characterized by high R&D expenditures, high barriers to entry, and high profit margins.

2. *Medical equipment industry* – The medical equipment industry is the second largest of the three industries, with sales of \$5.5 billion in 1997. It is characterized by high R&D expenditures, high barriers to entry, and high profit margins.

3. *Medical services industry* – The medical services industry is the third largest of the three industries, with sales of \$4.5 billion in 1997. It is characterized by high R&D expenditures, high barriers to entry, and high profit margins.

[illegible]

100

Figure 1 illustrates the experimental setup. A subject is seated at a table, viewing a video screen. A camera is positioned above the screen. A target is placed on the table. A horizontal arrow indicates the direction of movement. A vertical arrow indicates the direction of the video feedback. A horizontal arrow indicates the direction of the video feedback. A vertical arrow indicates the direction of the video feedback. A horizontal arrow indicates the direction of the video feedback. A vertical arrow indicates the direction of the video feedback.

- 2.2.1. 2.4.1
- 2.2.1. 4.4.4
- 2.2.2. 0.0.0
- 2.2.4. 0.0.0 0.0.0

$\frac{1}{2}$ $\frac{1}{2}$

424

<210> 250
<211> 430
<212> DNA
<213> Homo sapien

- 220. $\mathbf{A} = \begin{bmatrix} 1 & 2 \\ 2 & 1 \end{bmatrix}$
- 221. $\mathbf{A} = \begin{bmatrix} 1 & 2 & 3 \\ 2 & 1 & 4 \\ 3 & 4 & 1 \end{bmatrix}$
- 222. $\mathbf{A} = \begin{bmatrix} 1 & 2 & 3 & 4 \\ 2 & 1 & 4 & 3 \\ 3 & 4 & 1 & 2 \\ 4 & 3 & 2 & 1 \end{bmatrix}$
- 223. $\mathbf{A} = \begin{bmatrix} 1 & 2 & 3 \\ 2 & 1 & 4 \\ 3 & 4 & 1 \end{bmatrix}$

1000

Figure 1 illustrates the developmental stages of a child's drawing of a person. The sequence starts with a simple dot for a head at age 2 and gradually adds more features: a torso, limbs, and facial details like eyes, nose, and mouth, culminating in a more complex and realistic figure at age 10.

1. *Journal of the American Medical Association*, 1997; 277: 1033-1037.

1. *Chlorophyll a* (Chl *a*)

<220>

• 492 • 493

[illegible]

Figure 1: Schematic representation of the experimental design. The diagram shows a sequence of events: 'Stimulus presentation' (a box with a question mark), 'Response' (a box with a question mark), 'Feedback' (a box with a question mark), and 'Inter-trial interval' (a box with a question mark). The sequence is repeated for multiple trials, with a 'Block' label indicating a group of trials. The 'Inter-trial interval' is marked with a double slash '//'. The 'Stimulus presentation' box contains a question mark, and the 'Response' box contains a question mark. The 'Feedback' box contains a question mark. The 'Inter-trial interval' box contains a question mark. The sequence is repeated for multiple trials, with a 'Block' label indicating a group of trials. The 'Inter-trial interval' is marked with a double slash '//'. The 'Stimulus presentation' box contains a question mark, and the 'Response' box contains a question mark. The 'Feedback' box contains a question mark. The 'Inter-trial interval' box contains a question mark.

```

> 210> 239
> 211> 231
> 212> 194
> 213> Home region
> 221>
> 221> Miss feature
> 222> (1,..., 291)
> 223> n = A, T, C or G

```

- *211* 269
- *211* 438
- *214* FNA
- *213* Hinc earlier

1997, 1998, 1999, 2000, 2001, 2002, 2003, 2004, 2005, 2006, 2007, 2008, 2009, 2010, 2011, 2012, 2013, 2014, 2015, 2016, 2017, 2018, 2019, 2020, 2021, 2022, 2023, 2024, 2025, 2026, 2027, 2028, 2029, 2030, 2031, 2032, 2033, 2034, 2035, 2036, 2037, 2038, 2039, 2040, 2041, 2042, 2043, 2044, 2045, 2046, 2047, 2048, 2049, 2050, 2051, 2052, 2053, 2054, 2055, 2056, 2057, 2058, 2059, 2060, 2061, 2062, 2063, 2064, 2065, 2066, 2067, 2068, 2069, 2070, 2071, 2072, 2073, 2074, 2075, 2076, 2077, 2078, 2079, 2080, 2081, 2082, 2083, 2084, 2085, 2086, 2087, 2088, 2089, 2090, 2091, 2092, 2093, 2094, 2095, 2096, 2097, 2098, 2099, 2100, 2101, 2102, 2103, 2104, 2105, 2106, 2107, 2108, 2109, 2110, 2111, 2112, 2113, 2114, 2115, 2116, 2117, 2118, 2119, 2120, 2121, 2122, 2123, 2124, 2125, 2126, 2127, 2128, 2129, 2130, 2131, 2132, 2133, 2134, 2135, 2136, 2137, 2138, 2139, 2140, 2141, 2142, 2143, 2144, 2145, 2146, 2147, 2148, 2149, 2150, 2151, 2152, 2153, 2154, 2155, 2156, 2157, 2158, 2159, 2160, 2161, 2162, 2163, 2164, 2165, 2166, 2167, 2168, 2169, 2170, 2171, 2172, 2173, 2174, 2175, 2176, 2177, 2178, 2179, 2180, 2181, 2182, 2183, 2184, 2185, 2186, 2187, 2188, 2189, 2190, 2191, 2192, 2193, 2194, 2195, 2196, 2197, 2198, 2199, 2200, 2201, 2202, 2203, 2204, 2205, 2206, 2207, 2208, 2209, 2210, 2211, 2212, 2213, 2214, 2215, 2216, 2217, 2218, 2219, 2220, 2221, 2222, 2223, 2224, 2225, 2226, 2227, 2228, 2229, 2230, 2231, 2232, 2233, 2234, 2235, 2236, 2237, 2238, 2239, 2240, 2241, 2242, 2243, 2244, 2245, 2246, 2247, 2248, 2249, 2250, 2251, 2252, 2253, 2254, 2255, 2256, 2257, 2258, 2259, 2260, 2261, 2262, 2263, 2264, 2265, 2266, 2267, 2268, 2269, 2270, 2271, 2272, 2273, 2274, 2275, 2276, 2277, 2278, 2279, 2280, 2281, 2282, 2283, 2284, 2285, 2286, 2287, 2288, 2289, 2290, 2291, 2292, 2293, 2294, 2295, 2296, 2297, 2298, 2299, 2300, 2301, 2302, 2303, 2304, 2305, 2306, 2307, 2308, 2309, 2310, 2311, 2312, 2313, 2314, 2315, 2316, 2317, 2318, 2319, 2320, 2321, 2322, 2323, 2324, 2325, 2326, 2327, 2328, 2329, 2330, 2331, 2332, 2333, 2334, 2335, 2336, 2337, 2338, 2339, 2340, 2341, 2342, 2343, 2344, 2345, 2346, 2347, 2348, 2349, 2350, 2351, 2352, 2353, 2354, 2355, 2356, 2357, 2358, 2359, 2360, 2361, 2362, 2363, 2364, 2365, 2366, 2367, 2368, 2369, 2370, 2371, 2372, 2373, 2374, 2375, 2376, 2377, 2378, 2379, 2380, 2381, 2382, 2383, 2384, 2385, 2386, 2387, 2388, 2389, 2390, 2391, 2392, 2393, 2394, 2395, 2396, 2397, 2398, 2399, 2400, 2401, 2402, 2403, 2404, 2405, 2406, 2407, 2408, 2409, 2410, 2411, 2412, 2413, 2414, 2415, 2416, 2417, 2418, 2419, 2420, 2421, 2422, 2423, 2424, 2425, 2426, 2427, 2428, 2429, 2430, 2431, 2432, 2433, 2434, 2435, 2436, 2437, 2438, 2439, 2440, 2441, 2442, 2443, 2444, 2445, 2446, 2447, 2448, 2449, 2450, 2451, 2452, 2453, 2454, 2455, 2456, 2457, 2458, 2459, 2460, 2461, 2462, 2463, 2464, 2465, 2466, 2467, 2468, 2469, 2470, 2471, 2472, 2473, 2474, 2475, 2476, 2477, 2478, 2479, 2480, 2481, 2482, 2483, 2484, 2485, 2486, 2487, 2488, 2489, 2490, 2491, 2492, 2493, 2494, 2495, 2496, 2497, 2498, 2499, 2500, 2501, 2502, 2503, 2504, 2505, 2506, 2507, 2508, 2509, 2510, 2511, 2512, 2513, 2514, 2515, 2516, 2517, 2518, 2519, 2520, 2521, 2522, 2523, 2524, 2525, 2526, 2527, 2528, 2529, 2530, 2531, 2532, 2533, 2534, 2535, 2536, 2537, 2538, 2539, 2540, 2541, 2542, 2543, 2544, 2545, 2546, 2547, 2548, 2549, 2550, 2551, 2552, 2553, 2554, 2555, 2556, 2557, 2558, 2559, 2560, 2561, 2562, 2563, 2564, 2565, 2566, 2567, 2568, 2569, 2570, 2571, 2572, 2573, 2574, 2575, 2576, 2577, 2578, 2579, 2580, 2581, 2582, 2583, 2584, 2585, 2586, 2587, 2588, 2589, 2590, 2591, 2592, 2593, 2594, 2595, 2596, 2597, 2598, 2599, 2600, 2601, 2602, 2603, 2604, 2605, 2606, 2607, 2608, 2609, 2610, 2611, 2612, 2613, 2614, 2615, 2616, 2617, 2618, 2619, 2620, 2621, 2622, 2623, 2624, 2625, 2626, 2627, 2628, 2629, 2630, 2631, 2632, 2633, 2634, 2635, 2636, 2637, 2638, 2639, 2640, 2641, 2642, 2643, 2644, 2645, 2646, 2647, 2648, 2649, 2650, 2651, 2652, 2653, 2654, 2655, 2656, 2657, 2658, 2659, 2660, 2661, 2662, 2663, 2664, 2665, 2666, 2667, 2668, 2669, 2670, 2671, 2672, 2673, 2674, 2675, 2676, 2677, 2678, 26

the 1990s, the number of people in the world who are illiterate has increased from 1.2 billion to 1.5 billion. The number of illiterate people in the world is expected to reach 1.7 billion by the year 2015. The number of illiterate people in the world is expected to reach 1.7 billion by the year 2015. The number of illiterate people in the world is expected to reach 1.7 billion by the year 2015.

[illegible]

- 222.1. $\text{H}_2\text{O} \rightarrow \text{H}_2 + \frac{1}{2}\text{O}_2$
- 222.2. $\text{H}_2 + \frac{1}{2}\text{O}_2 \rightarrow \text{H}_2\text{O}$
- 222.3. $\text{H}_2 + \text{O}_2 \rightarrow \text{H}_2\text{O}_2$
- 222.4. $\text{H}_2 + \text{O}_2 \rightarrow \text{H}_2\text{O} + \text{O}$

| | | | | | | |
|------------|------------|-------------|-------------|-------------|------------|-----|
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| tttggttgtr | tctttttttt | cttttccttc | ccatatacctc | ctaattttaag | tttgacttgt | 120 |
| ttgtgagga | ggcaggagct | agagaactgt | gtgagctcat | aggggtggga | agttttactt | 180 |
| tcaagtcctg | cccactcact | actgtctctc | accttcctct | gaacaggctt | acaantgggt | 240 |
| tcttgctctg | tttccctttg | gacccaacaa | gcccctgtta | tgagtgtgca | tgactctgac | 300 |
| agctgtggac | ttagggctct | tggctacaga | tgcacgttaa | aatatctcat | ccagttcttg | 360 |
| caaatgttta | caataaacac | atttcctaga | ttccagtacc | caaatcatgt | ctttaaacac | 420 |
| tgtactctac | cccaagaagt | ggcacaataa | ctcttgggga | attatttaatt | tttttttttt | 480 |
| ctctctctac | gnnngnnng | gnnngnccag | gaattacaa | cttggaaagac | ctgggctgaa | 540 |
| ttttttatct | ctttttctct | attttttttt | ctaaacacaa | ccgtgttca | | 598 |

1. *Chlorophyll a* and *Chlorophyll b* were determined by the method of Arar and Collins (1971).
 2. *Chlorophyll a* and *Chlorophyll b* were determined by the method of Arar and Collins (1971).
 3. *Chlorophyll a* and *Chlorophyll b* were determined by the method of Arar and Collins (1971).

100

- * 2.1.1 - 204
- * 2.1.2 - 716
- * 2.1.3 - 738
- * 2.1.3 - Home Chapter

- 2.2.1. $\text{Info} = \text{feature}$
- 2.2.2. $\text{Info} = \text{feature} \cdot \text{Info}$

- *210* 265
- *211* 152
- *212* DNA
- *213* Home, earlier

| | | |
|-------|----------|-----|
| | Page 265 | |
| ***** | ***** | 80 |
| ***** | ***** | 120 |
| ***** | ***** | 180 |

| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
|---|---|---|---|---|---|---|---|---|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|----|-----|
| 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 | 13 | 14 | 15 | 16 | 17 | 18 | 19 | 20 | 21 | 22 | 23 | 24 | 25 | 26 | 27 | 28 | 29 | 30 | 31 | 32 | 33 | 34 | 35 | 36 | 37 | 38 | 39 | 40 | 41 | 42 | 43 | 44 | 45 | 46 | 47 | 48 | 49 | 50 | 51 | 52 | 53 | 54 | 55 | 56 | 57 | 58 | 59 | 60 | 61 | 62 | 63 | 64 | 65 | 66 | 67 | 68 | 69 | 70 | 71 | 72 | 73 | 74 | 75 | 76 | 77 | 78 | 79 | 80 | 81 | 82 | 83 | 84 | 85 | 86 | 87 | 88 | 89 | 90 | 91 | 92 | 93 | 94 | 95 | 96 | 97 | 98 | 99 | 100 |
| 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 | 13 | 14 | 15 | 16 | 17 | 18 | 19 | 20 | 21 | 22 | 23 | 24 | 25 | 26 | 27 | 28 | 29 | 30 | 31 | 32 | 33 | 34 | 35 | 36 | 37 | 38 | 39 | 40 | 41 | 42 | 43 | 44 | 45 | 46 | 47 | 48 | 49 | 50 | 51 | 52 | 53 | 54 | 55 | 56 | 57 | 58 | 59 | 60 | 61 | 62 | 63 | 64 | 65 | 66 | 67 | 68 | 69 | 70 | 71 | 72 | 73 | 74 | 75 | 76 | 77 | 78 | 79 | 80 | 81 | 82 | 83 | 84 | 85 | 86 | 87 | 88 | 89 | 90 | 91 | 92 | 93 | 94 | 95 | 96 | 97 | 98 | 99 | 100 |

<21> Homo sapiens

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<21> n = A,T,C or G

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100
100
100

<21> 100

<21> 100

<21> 100

<21> Homo sapiens

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ttctgaatg caatctatc agtaacaaa gtgtgttagg caatctatc gcatcttaa
ttcacaagaq cttataatga tctgggcttc tctcaggag gcaatcttc gagggaatg
gttcagagat gctcttcaa tctcagcttc aatctatct ctttggggg acacacataa
gtgtttttt gactcttcg atgcccagga gagagctctc aatctatga cttcattaga
aaaacactga atgtggggg caatctaaa attgttcagg aagcctctgt tcaagctgaa
tactggcatg aatctaaaa gaggatctg gatcgaaca

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400
420
440
460

<21> 100

<21> 100

<21> 100

<21> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(533)

<223> n = A,T,C or G

<401> 265

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aagtcggggg ggttcgatt taacgggaagc gctagctgca gtgggtctgc gcttcgggac
aaattctttg ggttgcttaa ggcggcgggg aatttgaggt atctctatca gtagtgagc
aagttgggac agtcgtaatt cccgaatccg attctcttga atctcagac cttcagagat
tgtctaatc atcaactga aggcagctat aattacggt tctgtctta ctagctccac
tcaataacta ggtgtctga cttctcttc gtagctga ggtcgttcg ttggaatttc
caactcgggt gatttgaga tttcaagttc cgaactgtt cgtctaacn atttgcatg
ttcagcatg aagcgaata aattcttca gtagcgggaa ttggatgga aca

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120
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240
300
360
420
480
540

<21> 100

<21> 100

<21> 100

<21> Homo sapiens

Figure 1. The effect of the number of trials on the number of correct responses. The number of correct responses was significantly higher than the number of incorrect responses in all cases. Error bars represent the standard error of the mean.

[illegible]

100

[illegible]

- * 2011年 2月
- * 2011年 4月
- * 2011年 6月
- * 2011年 8月

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1. $\frac{1}{2} \times \frac{1}{2} = \frac{1}{4}$
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The diagram illustrates the experimental setup. A participant is seated at a table, looking at a video screen. A camera is positioned above the screen. A target is placed on the table. A horizontal arrow indicates the direction of movement. A vertical arrow indicates the direction of the video feedback. A horizontal arrow indicates the direction of the video feedback. A vertical arrow indicates the direction of the video feedback. A horizontal arrow indicates the direction of the video feedback. A vertical arrow indicates the direction of the video feedback.

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Figure 1 illustrates the experimental setup. A subject is seated at a table, viewing a video screen. A camera is positioned above the screen. A target is placed on the table. A horizontal arrow indicates the direction of movement. A vertical arrow indicates the direction of the video feedback. A horizontal arrow indicates the direction of the video feedback. A vertical arrow indicates the direction of the video feedback. A horizontal arrow indicates the direction of the video feedback. A vertical arrow indicates the direction of the video feedback.

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the 1990s, the number of people in the world who are illiterate has increased from 1.2 billion to 1.5 billion. The number of illiterate people in the world is expected to reach 1.7 billion by the year 2015. The number of illiterate people in the world is expected to reach 1.7 billion by the year 2015. The number of illiterate people in the world is expected to reach 1.7 billion by the year 2015.

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 ttattttt ttattttt ttattttt ttattttt ttattttt ttattttt 900

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185 | 186 | 187 | 188 | 189 | 190 | 191 | 192 | 193 | 194 | 195 | 196 | 197 | 198 | 199 | 200 | 201 | 202 | 203 | 204 | 205 | 206 | 207 | 208 | 209 | 210 | 211 | 212 | 213 | 214 | 215 | 216 | 217 | 218 | 219 | 220 | 221 | 222 | 223 | 224 | 225 | 226 | 227 | 228 | 229 | 230 | 231 | 232 | 233 | 234 | 235 | 236 | 237 | 238 | 239 | 240 | 241 | 242 | 243 | 244 | 245 | 246 | 247 | 248 | 249 | 250 | 251 | 252 | 253 | 254 | 255 | 256 | 257 | 258 | 259 | 260 | 261 | 262 | 263 | 264 | 265 | 266 | 267 | 268 | 269 | 270 | 271 | 272 | 273 | 274 | 275 | 276 | 277 | 278 | 279 | 280 | 281 | 282 | 283 | 284 | 285 | 286 | 287 | 288 | 289 | 290 | 291 | 292 | 293 | 294 | 295 | 296 | 297 | 298 | 299 | 300 | 301 | 302 | 303 | 304 | 305 | 306 | 307 | 308 | 309 | 310 | 311 | 312 | 313 | 314 | 315 | 316 | 317 | 318 | 319 | 320 | 321 | 322 | 323 | 324 | 325 | 326 | 327 | 328 | 329 | 330 | 331 | 332 | 333 | 334 | 335 | 336 | 337 | 338 | 339 | 340 | 341 | 342 | 343 | 344 | 345 | 346 | 347 | 348 | 349 | 350 | 351 | 352 | 353 | 354 | 355 | 356 | 357 | 358 | 359 | 360 | 361 | 362 | 363 | 364 | 365 | 366 | 367 | 368 | 369 | 370 | 371 | 372 | 373 | 374 | 375 | 376 | 377 | 378 | 379 | 380 | 381 | 382 | 383 | 384 | 385 | 386 | 387 | 388 | 389 | 390 | 391 | 392 | 393 | 394 | 395 | 396 | 397 | 398 | 399 | 400 | 401 | 402 | 403 | 404 | 405 | 406 | 407 | 408 | 409 | 410 | 411 | 412 | 413 | 414 | 415 | 416 | 417 | 418 | 419 | 420 | 421 | 422 | 423 | 424 | 425 | 426 | 427 | 428 | 429 | 430 | 431 | 432 | 433 | 434 | 435 | 436 | 437 | 438 | 439 | 440 | 441 | 442 | 443 | 444 | 445 | 446 | 447 | 448 | 449 | 450 | 451 | 452 | 453 | 454 | 455 | 456 | 457 | 458 | 459 | 460 | 461 | 462 | 463 | 464 | 465 | 466 | 467 | 468 | 469 | 470 | 471 | 472 | 473 | 474 | 475 | 476 | 477 | 478 | 479 | 480 | 481 | 482 | 483 | 484 | 485 | 486 | 487 | 488 | 489 | 490 | 491 | 492 | 493 | 494 | 495 | 496 | 497 | 498 | 499 | 500 | 501 | 502 | 503 | 504 | 505 | 506 | 507 | 508 | 509 | 510 | 511 | 512 | 513 | 514 | 515 | 516 | 517 | 518 | 519 | 520 | 521 | 522 | 523 | 524 | 525 | 526 | 527 | 528 | 529 | 530 | 531 | 532 | 533 | 534 | 535 | 536 | 537 | 538 | 539 | 540 | 541 | 542 | 543 | 544 | 545 | 546 | 547 | 548 | 549 | 550 | 551 | 552 | 553 | 554 | 555 | 556 | 557 | 558 | 559 | 560 | 561 | 562 | 563 | 564 | 565 | 566 | 567 | 568 | 569 | 570 | 571 | 572 | 573 | 574 | 575 | 576 | 577 | 578 | 579 | 580 | 581 | 582 | 583 | 584 | 585 | 586 | 587 | 588 | 589 | 590 | 591 | 592 | 593 | 594 | 595 | 596 | 597 | 598 | 599 | 600 | 601 | 602 | 603 | 604 | 605 | 606 | 607 | 608 | 609 | 610 | 611 | 612 | 613 | 614 | 615 | 616 | 617 | 618 | 619 | 620 | 621 | 622 | 623 | 624 | 625 | 626 | 627 | 628 | 629 | 630 | 631 | 632 | 633 | 634 | 635 | 636 | 637 | 638 | 639 | 640 | 641 | 642 | 643 | 644 | 645 | 646 | 647 | 648 | 649 | 650 | 651 | 652 | 653 | 654 | 655 | 656 | 657 | 658 | 659 | 660 | 661 | 662 | 663 | 664 | 665 | 666 | 667 | 668 | 669 | 670 | 671 | 672 | 673 | 674 | 675 | 676 | 677 | 678 | 679 | 680 | 681 | 682 | 683 | 684 | 685 | 686 | 687 | 688 | 689 | 690 | 691 | 692 | 693 | 694 | 695 | 696 | 697 | 698 | 699 | 700 | 701 | 702 | 703 | 704 | 705 | 706 | 707 | 708 | 709 | 710 | 711 | 712 | 713 | 714 | 715 | 716 | 717 | 718 | 719 | 720 | 721 | 722 | 723 | 724 | 725 | 726 | 727 | 728 | 729 | 730 | 731 | 732 | 733 | 734 | 735 | 736 | 737 | 738 | 739 | 740 | 741 | 742 | 743 | 744 | 745 | 746 | 747 | 748 | 749 | 750 | 751 | 752 | 753 | 754 | 755 | 756 | 757 | 758 | 759 | 760 | 761 | 762 | 763 | 764 | 765 | 766 | 767 | 768 | 769 | 770 | 771 | 772 | 773 | 774 | 775 | 776 | 777 | 778 | 779 | 780 | 781 | 782 | 783 | 784 | 785 | 786 | 787 | 788 | 789 | 790 | 791 | 792 | 793 | 794 | 795 | 796 | 797 | 798 | 799 | 800 | 801 | 802 | 803 | 804 | 805 | 806 | 807 | 808 | 809 | 810 | 811 | 812 | 813 | 814 | 815 | 816 | 817 | 818 | 819 | 820 | 821 | 822 | 823 | 824 | 825 | 826 | 827 | 828 | 829 | 830 | 831 | 832 | 833 | 834 | 835 | 836 | 837 | 838 | 839 | 840 | 841 | 842 | 843 | 844 | 845 | 846 | 847 | 848 | 849 | 850 | 851 | 852 | 853 | 854 | 855 | 856 | 857 | 858 | 859 | 860 | 861 | 862 | 863 | 864 | 865 | 866 | 867 | 868 | 869 | 870 | 871 | 872 | 873 | 874 | 875 | 876 | 877 | 878 | 879 | 880 | 881 | 882 | 883 | 884 | 885 | 886 | 887 | 888 | 889 | 890 | 891 | 892 | 893 | 894 | 895 | 896 | 897 | 898 | 899 | 900 | 901 | 902 | 903 | 904 | 905 | 906 | 907 | 908 | 909 | 910 | 911 | 912 | 913 | 914 | 915 | 916 | 917 | 918 | 919 | 920 | 921 | 922 | 923 | 924 | 925 | 926 | 927 | 928 | 929 | 930 | 931 | 932 | 933 | 934 | 935 | 936 | 937 | 938 | 939 | 940 | 941 | 942 | 943 | 944 | 945 | 946 | 947 | 948 | 949 | 950 | 951 | 952 | 953 | 954 | 955 | 956 | 957 | 958 | 959 | 960 | 961 | 962 | 963 | 964 | 965 | 966 | 967 | 968 | 969 | 970 | 971 | 972 | 973 | 974 | 975 | 976 | 977 | 978 | 979 | 980 | 981 | 982 | 983 | 984 | 985 | 986 | 987 | 988 | 989 | 990 | 991 | 992 | 993 | 994 | 995 | 996 | 997 | 998 | 999 | 1000 | 1001 | 1002 | 1003 | 1004 | 1005 | 1006 | 1007 | 1008 | 1009 | 1010 | 1011 | 1012 | 1013 | 1014 | 1015 | 1016 | 1017 | 1018 | 1019 | 1020 | 1021 | 1022 | 1023 | 1024 | 1025 | 1026 | 1027 | 1028 | 1029 | 1030 | 1031 | 1032 | 1033 | 1034 | 1035 | 1036 | 1037 | 1038 | 1039 | 1040 | 1041 | 1042 | 1043 | 1044 | 1045 | 1046 | 1047 | 1048 | 1049 | 1050 | 1051 | 1052 | 1053 | 1054 | 1055 | 1056 | 1057 | 1058 | 1059 | 1060 | 1061 | 1062 | 1063 | 1064 | 1065 | 1066 | 1067 | 1068 | 1069 | 1070 | 1071 | 1072 | 1073 | 1074 | 1075 | 1076 | 1077 | 1078 | 1079 | 1080 | 1081 | 1082 | 1083 | 1084 | 1085 | 1086 | 1087 | 1088 | 1089 | 1090 | 1091 | 1092 | 1093 | 1094 | 1095 | 1096 | 1097 | 1098 | 1099 | 1100 | 1101 | 1102 | 1103 | 1104 | 1105 | 1106 | 1107 | 1108 | 1109 | 1110 | 1111 | 1112 | 1113 | 1114 | 1115 | 1116 | 1117 | 1118 | 1119 | 1120 | 1121 | 1122 | 1123 | 1124 | 1125 | 1126 | 1127 | 1128 | 1129 | 1130 | 1131 | 1132 | 1133 | 1134 | 1135 | 1136 | 1137 | 1138 | 1139 | 1140 | 1141 | 1142 | 1143 | 1144 | 1145 | 1146 | 1147 | 1148 | 1149 | 1150 | 1151 | 1152 | 1153 | 1154 | 1155 | 1156 | 1157 | 1158 | 1159 | 1160 | 1161 | 1162 | 1163 | 1164 | 1165 | 1166 | 1167 | 1168 | 1169 | 1170 | 1171 | 1172 | 1173 | 1174 | 1175 | 1176 | 1177 | 1178 | 1179 | 1180 | 1181 | 1182 | 1183 | 1184 | 1185 | 1186 | 1187 | 1188 | 1189 | 1190 | 1191 | 1192 | 1193 | 1194 | 1195 | 1196 | 1197 | 1198 | 1199 | 1200 | 1201 | 1202 | 1203 | 1204 | 1205 | 1206 | 1207 | 1208 | 1209 | 1210 | 1211 | 1212 | 1213 | 1214 | 1215 | 1216 | 1217 | 1218 | 1219 | 1220 | 1221 | 1222 | 1223 | 1224 | 1225 | 1226 | 1227 | 1228 | 1229 | 1230 | 1231 | 1232 | 1233 | 1234 | 1235 | 1236 | 1237 | 1238 | 1239 | 1240 | 1241 | 1242 | 1243 | 1244 | 1245 | 1246 | 1247 | 1248 | 1249 | 1250 | 1251 | 1252 | 1253 | 1254 | 1255 | 1256 | 1257 | 1258 | 1259 | 1260 | 1261 | 1262 | 1263 | 1264 | 1265 | 1266 | 1267 | 1268 | 1269 | 1270 | 1271 | 1272 | 1273 | 1274 | 1275 | 1276 | 1277 | 1278 | 1279 | 1280 | 1281 | 1282 | 1283 | 1284 | 1285 | 1286 | 1287 | 1288 | 1289 | 1290 | 1291 | 1292 | 1293 | 1294 | 1295 | 1296 | 1297 | 1298 | 1299 | 1300 | 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| Group | Trials | Correct Responses (Mean) |
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| No feedback | 1 | ~4.5 |
| | 2 | ~5.5 |
| | 3 | ~6.5 |
| Feedback | 1 | ~7.5 |
| | 2 | ~8.5 |
| | 3 | ~9.5 |

$\frac{1}{2} \times \frac{1}{2} = \frac{1}{4}$

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[illegible]

(210) 293

<211> 668

4212 • J. Neurosci., September 24, 2008 • 28(39):4205–4212

• 213 • Homo sapien.

4 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80 81 82 83 84 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100 101 102 103 104 105 106 107 108 109 110 111 112 113 114 115 116 117 118 119 120 121 122 123 124 125 126 127 128 129 130 131 132 133 134 135 136 137 138 139 140 141 142 143 144 145 146 147 148 149 150 151 152 153 154 155 156 157 158 159 160 161 162 163 164 165 166 167 168 169 170 171 172 173 174 175 176 177 178 179 180 181 182 183 184 185 186 187 188 189 190 191 192 193 194 195 196 197 198 199 200 201 202 203 204 205 206 207 208 209 210 211 212 213 214 215 216 217 218 219 220 221 222 223 224 225 226 227 228 229 230 231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252 253 254 255 256 257 258 259 260 261 262 263 264 265 266 267 268 269 270 271 272 273 274 275 276 277 278 279 280 281 282 283 284 285 286 287 288 289 290 291 292 293 294 295 296 297 298 299 300 301 302 303 304 305 306 307 308 309 310 311 312 313 314 315 316 317 318 319 320 321 322 323 324 325 326 327 328 329 330 331 332 333 334 335 336 337 338 339 340 341 342 343 344 345 346 347 348 349 350 351 352 353 354 355 356 357 358 359 360 361 362 363 364 365 366 367 368 369 370 371 372 373 374 375 376 377 378 379 380 381 382 383 384 385 386 387 388 389 390 391 392 393 394 395 396 397 398 399 400 401 402 403 404 405 406 407 408 409 410 411 412 413 414 415 416 417 418 419 420 421 422 423 424 425 426 427 428 429 430 431 432 433 434 435 436 437 438 439 440 441 442 443 444 445 446 447 448 449 450 451 452 453 454 455 456 457 458 459 460 461 462 463 464 465 466 467 468 469 470 471 472 473 474 475 476 477 478 479 480 481 482 483 484 485 486 487 488 489 490 491 492 493 494 495 496 497 498 499 500 501 502 503 504 505 506 507 508 509 510 511 512 513 514 515 516 517 518 519 520 521 522 523 524 525 526 527 528 529 530 531 532 533 534 535 536 537 538 539 540 541 542 543 544 545 546 547 548 549 550 551 552 553 554 555 556 557 558 559 560 561 562 563 564 565 566 567 568 569 570 571 572 573 574 575 576 577 578 579 580 581 582 583 584 585 586 587 588 589 590 591 592 593 594 595 596 597 598 599 600 601 602 603 604 605 606 607 608 609 610 611 612 613 614 615 616 617 618 619 620 621 622 623 624 625 626 627 628 629 630 631 632 633 634 635 636 637 638 639 640 641 642 643 644 645 646 647 648 649 650 651 652 653 654 655 656 657 658 659 660 661 662 663 664 665 666 667 668 669 670 671 672 673 674 675 676 677 678 679 680 681 682 683 684 685 686 687 688 689 690 691 692 693 694 695 696 697 698 699 700 701 702 703 704 705 706 707 708 709 710 711 712 713 714 715 716 717 718 719 720 721 722 723 724 725 726 727 728 729 730 731 732 733 734 735 736 737 738 739 740 741 742 743 744 745 746 747 748 749 750 751 752 753 754 755 756 757 758 759 760 761 762 763 764 765 766 767 768 769 770 771 772 773 774 775 776 777 778 779 780 781 782 783 784 785 786 787 788 789 790 791 792 793 794 795 796 797 798 799 800 801 802 803 804 805 806 807 808 809 810 811 812 813 814 815 816 817 818 819 820 821 822 823 824 825 826 827 828 829 830 831 832 833 834 835 836 837 838 839 840 841 842 843 844 845 846 847 848 849 850 851 852 853 854 855 856 857 858 859 860 861 862 863 864 865 866 867 868 869 870 871 872 873 874 875 876 877 878 879 880 881 882 883 884 885 886 887 888 889 890 891 892 893 894 895 896 897 898 899 900 901 902 903 904 905 906 907 908 909 910 911 912 913 914 915 916 917 918 919 920 921 922 923 924 925 926 927 928 929 930 931 932 933 934 935 936 937 938 939 940 941 942 943 944 945 946 947 948 949 950 951 952 953 954 955 956 957 958 959 960 961 962 963 964 965 966 967 968 969 970 971 972 973 974 975 976 977 978 979 980 981 982 983 984 985 986 987 988 989 990 991 992 993 994 995 996 997 998 999 1000 1001 1002 1003 1004 1005 1006 1007 1008 1009 1010 1011 1012 1013 1014 1015 1016 1017 1018 1019 1020 1021 1022 1023 1024 1025 1026 1027 1028 1029 1030 1031 1032 1033 1034 1035 1036 1037 1038 1039 1040

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| ttcaaaacag | ttggaaaac | ggagttact | gtgagttgt | gaaactggt | ggtagacccc |
| atgtgttggg | taatactgg | ttctcttgg | tgtaaaaag | agatgggtg | tgaggttgat |
| tccatgggg | gtggttttt | gttgaagag | ccatttggc | tccaggagaa | gatggggaag |
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| ggagacagc | agaaactct | tatgaagaa | ctcaggagaa | agatgggaa | atgtgtggg |
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| caaaatgtt | agagcaaat | gaagatctt | ctaatcttg | cttgagga | tggaattca |
| gggtgtttt | gaatgtgtt | gtgaatctt | gtgaatctt | gttgagga | tggaattca |
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Figure 1 is a line graph showing the percentage of total protein in the supernatant fraction of the Golgi apparatus for various proteins. The y-axis is labeled 'PERCENTAGE OF TOTAL PROTEIN IN SUPERNATANT FRACTION OF GOLGI APPARATUS' and ranges from 0 to 100. The x-axis is labeled 'PROTEIN' and lists various proteins: α , β , γ , δ , ϵ , ζ , η , θ , ι , κ , λ , μ , ν , ξ , \omicron , π , ρ , σ , τ , ϕ , χ , ψ , ω , and others. The graph shows that most proteins are concentrated in the supernatant fraction, with some showing higher percentages in the pellet fraction.

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213- Homo sapien.

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Met Asp Ile Val Val Ser Gly Ser His Pro Leu Thr Val Asp Ser Phe
 1 5 15 15
 Leu His Leu Ala Gly Ser Asp Leu Leu Ser Arg Ser Leu Met Ala Glu
 20 25 30
 Glu Tyr Thr Ile Val His Ala Ser Phe Ile Ser Cys Ile Ser Ser Ser
 35 40 45
 Leu Asp Gly Glu Gly Glu Arg Glu Glu Gln Arg Gly His Phe Trp Arg
 50 55 60
 Pro Gln Arg Leu Leu Cys Glu Asp Ala Trp Glu Gln Glu Val Gln Val
 65 70 75 80
 Val Leu Pro Leu Leu Pro Leu Leu Gln Gly Ser Gly Lys Ser Asn Val
 85 90 95
 Val Ala Trp Gly Asp Tyr Asp Asp Ser Ala Phe Met Asp Pro Arg Tyr
 100 105 110
 His Val His Gly Glu Asp Leu Asp Lys Leu His Arg Ala Ala Trp Trp
 115 120 125
 Gly Lys Val Pro Arg Lys Asp Leu Ile Val Met Leu Arg Asp Thr Asp
 130 135 140
 Val Asn Lys Arg Asp Lys Gln Lys Arg Thr Ala Leu His Leu Ala Ser
 145 150 155 16
 Ala Asn Gly Asn Ser Glu Val Val Lys Leu Val Leu Asp Arg Arg Cys
 165 170 175
 Glu Leu Asn Val Leu Asp Asn Lys Lys Arg Thr Ala Leu Thr Lys Ala
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 His His Gly Leu Ala Ser Lys Met Gly Lys Thr Lys Lys Ala Lys His
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 His Lys Lys Ala His Ser Gly Lys Ser Asn Val Gly Thr Ser Gly Asp
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 Arg Ala Ala Thr Thr Gly Lys Val Pro Arg Lys Asp Leu Ile Val Met
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 Thr Thr Leu His Tyr Ala Ile Tyr Asn Gln Asp Lys Leu Met Ala Lys
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 Leu Thr Pro Leu Leu Leu Gly Val His Glu Gln Lys Gln Gln Val Val
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 Lys Phe Leu Ile Lys Lys Lys Ala Asn Leu Asn Ala Leu Asp Arg Tyr
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 Gly Arg Thr Ala Leu Ile Leu Ala Val Cys Cys Gly Ser Ala Ser Ile
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 Val Ser Leu Leu Leu Gln Gln Asn Ile Asp Val Ser Ser Gln Asp Leu
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 Ser Gly Gln Thr Ala Arg His Tyr Ala Val Ser Ser His His His Val
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 Ile Cys His Leu Leu Ser Asp Tyr Lys Gln Lys Gln Met Leu Lys Ile
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 Ser Ser Gln Asn Ser Asn Pro His Asn Val Ser Arg Thr Arg Asn Lys
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Met Val Thr Ala Val Asp Ser Met His Ala Ala Ser Ser Val Lys Lys
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 12 Lys Lys Arg Glu Ser Gly Lys Ser Asn Val Gly Thr Ser Gly Asp
 3
 His Asp Asp Ser Ala Met Lys Thr Leu Arg Ser Lys Met Gly Lys Thr
 4
 Lys Arg His Lys Thr His Lys Lys Arg Gly Ser Gly Lys Ser Asn Val
 5
 Gly Ala Ser Gly Asp His Asp Asp Ser Ala Met Lys Thr Leu Arg Asn
 6
 Lys Met Gly Lys Thr Lys Lys His Lys Thr His Lys Lys Arg Gly Ser
 7
 Gly Lys Ser Lys Val Gly Ala Thr Gly Asp Tyr Asp Asp Ser Ala Thr
 8
 Met Ala His Arg Tyr His Val Arg Gly Glu Asp Leu Asp Lys Leu His
 9
 Arg Ala Ala Tyr Tyr Gly Lys Val Pro Arg Lys Asp Leu Ile Val Met
 10
 Leu Arg Asp Thr Asp Val Asn Lys Lys Asp Lys Glu Lys Arg Thr Ala
 11
 Leu His Leu Ala Ser Ala Asn Gly Asn Ser Glu Val Val Lys Leu Leu
 12
 Leu Asp Arg Arg Cys Glu Leu Asn Val Leu Asp Asn Lys Lys Arg Thr
 13
 Ala Leu Ile Lys Ala Val Glu Cys Glu Glu Asp Glu Cys Ala Leu Met
 14
 Leu Leu His His Gly Thr Asp Pro Asn Ile His Asp Glu Tyr Gly Asn
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 Thr Thr Leu His Tyr Ala Ile Tyr Asn Glu Asp Lys Leu Met Ala Lys
 16
 Ala Leu Leu Leu Tyr Gly Ala Asp Ile Glu Ser Lys Asn Lys His Gly
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 Leu Thr Pro Leu Leu Leu Gly Val His Glu Glu Lys Glu Glu Val Val
 18
 Lys Phe Leu Ile Lys Lys Lys Ala Asn Leu Asn Ala Leu Asp Arg Tyr
 19
 Gly Arg Thr Ala Leu Ile Leu Ala Val Cys Cys Gly Ser Ala Ser Ile
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 Val Ser Leu Leu Leu Glu Glu Asn Ile Asp Val Ser Ser Glu Asp Leu
 21
 Ser Gly Glu Thr Ala Arg Glu Tyr Ala Val Ser Ser His His His Val
 22
 Ile Cys Glu Leu Leu Ser Asp Tyr Lys Glu Lys Glu Met Leu Lys Ile
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 Ser Ser His Asn Ser Asn Pro Glu Glu Asp Leu Lys Leu Thr Ser Glu
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 Glu Glu Ser His Arg His Lys Gly Ser His Asn Ser His His His Lys
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 Met Ser His His His His Ile Asn Lys Asp Gly Asp Arg His Val His
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 53 53 53 53
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 His Asp Asp Ser Ala Met Lys Thr Leu Arg Ser Lys Met Gly Lys Trp
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 Cys Arg His Cys Phe Pro Cys Cys Arg Gly Ser Gly Lys Ser Asn Val
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 Thr Thr Leu His Tyr Ala Leu Tyr Asn Glu Asp Lys Leu Met Ala Lys
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 Ala Leu Leu Leu Tyr Gly Ala Asp Ile Glu Ser Lys Asn Lys His Gly
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 Leu Thr Leu Leu Leu Leu Gly Val His Glu Glu Lys Glu Glu Val Val
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 Lys His Leu Ile Lys Lys Lys Ala Asn Leu Asn Ala Leu Asp Arg Tyr
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 Gly Arg Thr Ala Leu Ile Leu Ala Val Cys Cys Gly Ser Ala Ser Ile
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 Val Ser Leu Leu Leu Glu Glu Asn Ile Asp Val Ser Ser Glu Asp Leu
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 Glu Glu Met Lys Lys His Glu Ser Asn Asn Val Gly Leu Leu Glu Asn
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| Ser | Pro | Ileu | Lys | Gly | Ile | Ileu | Ser | Gln | Trp | Asp | Lys | Phe | Asp | Pr | Gln |
| | | | 20 | | | | | 23 | | | | | 30 | | |
| Thr | Ileu | Gln | Lys | Gln | Val | Ala | His | Phe | Phe | Gly | Thr | Met | Ala | Trp | Pro |
| | 24 | | | | | | 4 | | | | | 47 | | | |
| Gln | His | Ser | Ileu | Ser | Asp | Gly | Gln | Lys | Trp | Pr | Pr | Gln | Gly | Ser | Thr |
| | 51 | | | | | 54 | | | | | | 61 | | | |

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Lys Lys Thr Asn Ile Thr Lys Val Ala Ala Ile Ile Thr Leu Lys Ala

Asn Thr Leu Tyr Lys Ala

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Lys Met Ala Lys Ala Leu Leu Leu

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<400> 11

Gly Leu Thr Pro Leu Leu Gly Ile

<210> 12
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<400> 12

Lys Leu Val Leu Arg Arg Arg Tyr Glu Leu

the 1990s, the number of people in the world who are illiterate has increased from 1.2 billion to 1.5 billion. The number of illiterate people in the world is expected to reach 1.7 billion by the year 2015. The number of illiterate people in the world is expected to reach 1.7 billion by the year 2015. The number of illiterate people in the world is expected to reach 1.7 billion by the year 2015.

[illegible]

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- 211 • 679
- 212 • DNA
- 213 • Homo sapiens

[illegible]

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1. *Journal of the American Medical Association*, 1997; 277: 1033-1038.

1999, 2000, 2001, 2002, 2003, 2004, 2005, 2006, 2007, 2008, 2009, 2010, 2011, 2012, 2013, 2014, 2015, 2016, 2017, 2018, 2019, 2020, 2021, 2022, 2023, 2024, 2025, 2026, 2027, 2028, 2029, 2030, 2031, 2032, 2033, 2034, 2035, 2036, 2037, 2038, 2039, 2040, 2041, 2042, 2043, 2044, 2045, 2046, 2047, 2048, 2049, 2050, 2051, 2052, 2053, 2054, 2055, 2056, 2057, 2058, 2059, 2060, 2061, 2062, 2063, 2064, 2065, 2066, 2067, 2068, 2069, 2070, 2071, 2072, 2073, 2074, 2075, 2076, 2077, 2078, 2079, 2080, 2081, 2082, 2083, 2084, 2085, 2086, 2087, 2088, 2089, 2090, 2091, 2092, 2093, 2094, 2095, 2096, 2097, 2098, 2099, 2100, 2101, 2102, 2103, 2104, 2105, 2106, 2107, 2108, 2109, 2110, 2111, 2112, 2113, 2114, 2115, 2116, 2117, 2118, 2119, 2120, 2121, 2122, 2123, 2124, 2125, 2126, 2127, 2128, 2129, 2130, 2131, 2132, 2133, 2134, 2135, 2136, 2137, 2138, 2139, 2140, 2141, 2142, 2143, 2144, 2145, 2146, 2147, 2148, 2149, 2150, 2151, 2152, 2153, 2154, 2155, 2156, 2157, 2158, 2159, 2160, 2161, 2162, 2163, 2164, 2165, 2166, 2167, 2168, 2169, 2170, 2171, 2172, 2173, 2174, 2175, 2176, 2177, 2178, 2179, 2180, 2181, 2182, 2183, 2184, 2185, 2186, 2187, 2188, 2189, 2190, 2191, 2192, 2193, 2194, 2195, 2196, 2197, 2198, 2199, 2200, 2201, 2202, 2203, 2204, 2205, 2206, 2207, 2208, 2209, 2210, 2211, 2212, 2213, 2214, 2215, 2216, 2217, 2218, 2219, 2220, 2221, 2222, 2223, 2224, 2225, 2226, 2227, 2228, 2229, 2230, 2231, 2232, 2233, 2234, 2235, 2236, 2237, 2238, 2239, 2240, 2241, 2242, 2243, 2244, 2245, 2246, 2247, 2248, 2249, 2250, 2251, 2252, 2253, 2254, 2255, 2256, 2257, 2258, 2259, 2260, 2261, 2262, 2263, 2264, 2265, 2266, 2267, 2268, 2269, 2270, 2271, 2272, 2273, 2274, 2275, 2276, 2277, 2278, 2279, 2280, 2281, 2282, 2283, 2284, 2285, 2286, 2287, 2288, 2289, 2290, 2291, 2292, 2293, 2294, 2295, 2296, 2297, 2298, 2299, 2300, 2301, 2302, 2303, 2304, 2305, 2306, 2307, 2308, 2309, 2310, 2311, 2312, 2313, 2314, 2315, 2316, 2317, 2318, 2319, 2320, 2321, 2322, 2323, 2324, 2325, 2326, 2327, 2328, 2329, 2330, 2331, 2332, 2333, 2334, 2335, 2336, 2337, 2338, 2339, 2340, 2341, 2342, 2343, 2344, 2345, 2346, 2347, 2348, 2349, 2350, 2351, 2352, 2353, 2354, 2355, 2356, 2357, 2358, 2359, 2360, 2361, 2362, 2363, 2364, 2365, 2366, 2367, 2368, 2369, 2370, 2371, 2372, 2373, 2374, 2375, 2376, 2377, 2378, 2379, 2380, 2381, 2382, 2383, 2384, 2385, 2386, 2387, 2388, 2389, 2390, 2391, 2392, 2393, 2394, 2395, 2396, 2397, 2398, 2399, 2400, 2401, 2402, 2403, 2404, 2405, 2406, 2407, 2408, 2409, 2410, 2411, 2412, 2413, 2414, 2415, 2416, 2417, 2418, 2419, 2420, 2421, 2422, 2423, 2424, 2425, 2426, 2427, 2428, 2429, 2430, 2431, 2432, 2433, 2434, 2435, 2436, 2437, 2438, 2439, 2440, 2441, 2442, 2443, 2444, 2445, 2446, 2447, 2448, 2449, 2450, 2451, 2452, 2453, 2454, 2455, 2456, 2457, 2458, 2459, 2460, 2461, 2462, 2463, 2464, 2465, 2466, 2467, 2468, 2469, 2470, 2471, 2472, 2473, 2474, 2475, 2476, 2477, 2478, 2479, 2480, 2481, 2482, 2483, 2484, 2485, 2486, 2487, 2488, 2489, 2490, 2491, 2492, 2493, 2494, 2495, 2496, 2497, 2498, 2499, 2500, 2501, 2502, 2503, 2504, 2505, 2506, 2507, 2508, 2509, 2510, 2511, 2512, 2513, 2514, 2515, 2516, 2517, 2518, 2519, 2520, 2521, 2522, 2523, 2524, 2525, 2526, 2527, 2528, 2529, 2530, 2531, 2532, 2533, 2534, 2535, 2536, 2537, 2538, 2539, 2540, 2541, 2542, 2543, 2544, 2545, 2546, 2547, 2548, 2549, 2550, 2551, 2552, 2553, 2554, 2555, 2556, 2557, 2558, 2559, 2560, 2561, 2562, 2563, 2564, 2565, 2566, 2567, 2568, 2569, 2570, 2571, 2572, 2573, 2574, 2575, 2576, 2577, 2578, 2579, 2580, 2581, 2582, 2583, 2584, 2585, 2586, 2587, 2588, 2589, 2590, 2591, 2592, 2593, 2594, 2595, 2596, 2597, 2598, 2599, 2600, 2601, 2602, 2603, 2604, 2605, 2606, 2607, 2608, 2609, 2610, 2611, 2612, 2613, 2614, 2615, 2616, 2617, 2618, 2619, 2620, 2621, 2622, 2623, 2624, 2625, 2626, 2627, 2628, 2629, 2630, 2631, 2632, 2633, 2634, 2635, 2636, 2637, 2638, 2639, 2640, 2641, 2642, 2643, 2644, 2645, 2646, 2647, 2648, 2649, 2650, 2651, 2652, 2653, 2654, 2655, 2656, 2657, 2658, 2659, 2660, 2661, 2662, 2663, 2664, 2665, 2666, 2667, 2668, 2669, 2670, 2671, 2672, 2673, 2674, 2675, 2676, 2677, 2678, 2679, 2680, 26

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| aaagatgag | aaatagag | aaagagag | atgagatag | aaatagat | aaatgagat | 14 |
| aaagttata | taaggtat | agttttat | agtttgggt | ttatgaaa | aaagggat | 16 |
| gaaagag | taaaatgt | gttgggag | gttgggtg | aaatgtat | atgtat | 24 |
| gaaagatga | taaggggt | agaggggt | agatgaat | gttgaaggt | atggggat | 30 |
| aggttaag | ggaatgat | aggtgaggt | atgtatgga | gttgggaat | aaatgggt | 36 |
| ggaggggta | gaggggaagt | gagatgggt | gaggggaat | gttgggaat | aaatgggt | 42 |
| aaatgggga | ggttatgt | gttgggtt | gattgaatg | gaggtttt | ttttggaat | 48 |
| aaatgatat | gtatgaggt | aaatgggt | aaatgtat | ggtttgt | aaatggtt | 54 |
| aaagagatg | gaaagata | gttgggtt | gttgggaat | agggggt | ttttatggt | 60 |
| aaatgaata | aaagatgt | aaatgggt | gaaaggtgt | gagggggt | gagggggt | 66 |
| aaagagatg | aaatgtgt | gttgggtt | gaaaggt | ggttatgaa | ggttatgaa | 72 |
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| gttggggaag | atgtggaaa | gttgaaga | gttgggtgt | gttgggtg | gttgaagaat | 90 |
| gatgtcatg | ttatgtcat | gttgaatgaa | gttgaaga | gttgaaga | aaatgggt | 96 |
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| gttgtcat | gttgtatg | gttgaaga | gttgaaga | gttgaaga | gttgaaga | 132 |
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| gttgtcat | gttgtatg | gttgaaga | gttgaaga | gttgaaga | gttgaaga | 300 |
| gttgtcat | gttgtatg | gttgaaga | gttgaaga | gttgaaga | gttgaaga | 306 |
| gttgtcat | gttgtatg | gttgaaga | gttgaaga | gttgaaga | gttgaaga | 312 |
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| gttgtcat | gttgtatg | gttgaaga | gttgaaga | gttgaaga | gttgaaga | 324 |
| gttgtcat | gttgtatg | gttgaaga | gttgaaga | gttgaaga | gttgaaga | 330 |
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| gttgtcat | gttgtatg | gttgaaga | gttgaaga | gttgaaga | gttgaaga | 354 |
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|---|---|---|---|
| 1. $\frac{1}{2} \times \frac{1}{3} = \frac{1}{6}$ | 2. $\frac{1}{4} \times \frac{1}{5} = \frac{1}{20}$ | 3. $\frac{1}{6} \times \frac{1}{7} = \frac{1}{42}$ | 4. $\frac{1}{8} \times \frac{1}{9} = \frac{1}{72}$ |
| 5. $\frac{1}{10} \times \frac{1}{11} = \frac{1}{110}$ | 6. $\frac{1}{12} \times \frac{1}{13} = \frac{1}{156}$ | 7. $\frac{1}{14} \times \frac{1}{15} = \frac{1}{210}$ | 8. $\frac{1}{16} \times \frac{1}{17} = \frac{1}{272}$ |
| 9. $\frac{1}{18} \times \frac{1}{19} = \frac{1}{342}$ | 10. $\frac{1}{20} \times \frac{1}{21} = \frac{1}{420}$ | 11. $\frac{1}{22} \times \frac{1}{23} = \frac{1}{506}$ | 12. $\frac{1}{24} \times \frac{1}{25} = \frac{1}{600}$ |
| 13. $\frac{1}{26} \times \frac{1}{27} = \frac{1}{702}$ | 14. $\frac{1}{28} \times \frac{1}{29} = \frac{1}{812}$ | 15. $\frac{1}{30} \times \frac{1}{31} = \frac{1}{930}$ | 16. $\frac{1}{32} \times \frac{1}{33} = \frac{1}{1056}$ |
| 17. $\frac{1}{34} \times \frac{1}{35} = \frac{1}{1190}$ | 18. $\frac{1}{36} \times \frac{1}{37} = \frac{1}{1332}$ | 19. $\frac{1}{38} \times \frac{1}{39} = \frac{1}{1482}$ | 20. $\frac{1}{40} \times \frac{1}{41} = \frac{1}{1640}$ |

21. $\frac{1}{42} \times \frac{1}{43} = \frac{1}{1806}$

22. $\frac{1}{44} \times \frac{1}{45} = \frac{1}{1980}$

23. $\frac{1}{46} \times \frac{1}{47} = \frac{1}{2162}$

24. $\frac{1}{48} \times \frac{1}{49} = \frac{1}{2352}$

25. $\frac{1}{50} \times \frac{1}{51} = \frac{1}{2550}$

26. $\frac{1}{52} \times \frac{1}{53} = \frac{1}{2756}$

27. $\frac{1}{54} \times \frac{1}{55} = \frac{1}{2970}$

28. $\frac{1}{56} \times \frac{1}{57} = \frac{1}{3192}$

29. $\frac{1}{58} \times \frac{1}{59} = \frac{1}{3422}$

30. $\frac{1}{60} \times \frac{1}{61} = \frac{1}{3660}$

31. $\frac{1}{62} \times \frac{1}{63} = \frac{1}{3906}$

32. $\frac{1}{64} \times \frac{1}{65} = \frac{1}{4160}$

33. $\frac{1}{66} \times \frac{1}{67} = \frac{1}{4422}$

34. $\frac{1}{68} \times \frac{1}{69} = \frac{1}{4692}$

35. $\frac{1}{70} \times \frac{1}{71} = \frac{1}{4970}$

36. $\frac{1}{72} \times \frac{1}{73} = \frac{1}{5256}$

37. $\frac{1}{74} \times \frac{1}{75} = \frac{1}{5550}$

38. $\frac{1}{76} \times \frac{1}{77} = \frac{1}{5852}$

39. $\frac{1}{78} \times \frac{1}{79} = \frac{1}{6162}$

40. $\frac{1}{80} \times \frac{1}{81} = \frac{1}{6480}$

41. $\frac{1}{82} \times \frac{1}{83} = \frac{1}{6806}$

42. $\frac{1}{84} \times \frac{1}{85} = \frac{1}{7140}$

43. $\frac{1}{86} \times \frac{1}{87} = \frac{1}{7482}$

44. $\frac{1}{88} \times \frac{1}{89} = \frac{1}{7840}$

45. $\frac{1}{90} \times \frac{1}{91} = \frac{1}{8190}$

46. $\frac{1}{92} \times \frac{1}{93} = \frac{1}{8556}$

47. $\frac{1}{94} \times \frac{1}{95} = \frac{1}{8930}$

48. $\frac{1}{96} \times \frac{1}{97} = \frac{1}{9312}$

49. $\frac{1}{98} \times \frac{1}{99} = \frac{1}{9702}$

50. $\frac{1}{100} \times \frac{1}{101} = \frac{1}{10100}$

51. $\frac{1}{102} \times \frac{1}{103} = \frac{1}{10506}$

52. $\frac{1}{104} \times \frac{1}{105} = \frac{1}{10920}$

53. $\frac{1}{106} \times \frac{1}{107} = \frac{1}{11342}$

54. $\frac{1}{108} \times \frac{1}{109} = \frac{1}{11772}$

55. $\frac{1}{110} \times \frac{1}{111} = \frac{1}{12210}$

56. $\frac{1}{112} \times \frac{1}{113} = \frac{1}{12656}$

57. $\frac{1}{114} \times \frac{1}{115} = \frac{1}{13110}$

58. $\frac{1}{116} \times \frac{1}{117} = \frac{1}{13572}$

59. $\frac{1}{118} \times \frac{1}{119} = \frac{1}{14042}$

60. $\frac{1}{120} \times \frac{1}{121} = \frac{1}{14520}$

61. $\frac{1}{122} \times \frac{1}{123} = \frac{1}{15006}$

62. $\frac{1}{124} \times \frac{1}{125} = \frac{1}{15500}$

63. $\frac{1}{126} \times \frac{1}{127} = \frac{1}{15992}$

64. $\frac{1}{128} \times \frac{1}{129} = \frac{1}{16480}$

65. $\frac{1}{130} \times \frac{1}{131} = \frac{1}{16970}$

66. $\frac{1}{132} \times \frac{1}{133} = \frac{1}{17462}$

67. $\frac{1}{134} \times \frac{1}{135} = \frac{1}{17952}$

68. $\frac{1}{136} \times \frac{1}{137} = \frac{1}{18440}$

69. $\frac{1}{138} \times \frac{1}{139} = \frac{1}{18922}$

70. $\frac{1}{140} \times \frac{1}{141} = \frac{1}{19400}$

71. $\frac{1}{142} \times \frac{1}{143} = \frac{1}{19872}$

72. $\frac{1}{144} \times \frac{1}{145} = \frac{1}{20340}$

73. $\frac{1}{146} \times \frac{1}{147} = \frac{1}{20802}$

74. $\frac{1}{148} \times \frac{1}{149} = \frac{1}{21260}$

75. $\frac{1}{150} \times \frac{1}{151} = \frac{1}{21710}$

76. $\frac{1}{152} \times \frac{1}{153} = \frac{1}{22162}$

77. $\frac{1}{154} \times \frac{1}{155} = \frac{1}{22610}$

78. $\frac{1}{156} \times \frac{1}{157} = \frac{1}{23052}$

79. $\frac{1}{158} \times \frac{1}{159} = \frac{1}{23490}$

80. $\frac{1}{160} \times \frac{1}{161} = \frac{1}{23920}$

81. $\frac{1}{162} \times \frac{1}{163} = \frac{1}{24352}$

82. $\frac{1}{164} \times \frac{1}{165} = \frac{1}{24780}$

83. $\frac{1}{166} \times \frac{1}{167} = \frac{1}{25202}$

84. $\frac{1}{168} \times \frac{1}{169} = \frac{1}{25620}$

85. $\frac{1}{170} \times \frac{1}{171} = \frac{1}{26030}$

86. $\frac{1}{172} \times \frac{1}{173} = \frac{1}{26442}$

87. $\frac{1}{174} \times \frac{1}{175} = \frac{1}{26850}$

88. $\frac{1}{176} \times \frac{1}{177} = \frac{1}{27252}$

89. $\frac{1}{178} \times \frac{1}{179} = \frac{1}{27650}$

90. $\frac{1}{180} \times \frac{1}{181} = \frac{1}{28040}$

91. $\frac{1}{182} \times \frac{1}{183} = \frac{1}{28432}$

92. $\frac{1}{184} \times \frac{1}{185} = \frac{1}{28820}$

93. $\frac{1}{186} \times \frac{1}{187} = \frac{1}{29202}$

94. $\frac{1}{188} \times \frac{1}{189} = \frac{1}{29580}$

95. $\frac{1}{190} \times \frac{1}{191} = \frac{1}{29950}$

96. $\frac{1}{192} \times \frac{1}{193} = \frac{1}{30322}$

97. $\frac{1$

1. *Journal of the American Medical Association*, 1997; 277: 1033-1037.

Met Ser Asp Asp Thr Asp Val Asn Lys Lys Asp Lys Glu Lys Asp Thr
411 412 413 414 415 416 417 418 419 420

Ala Leu His Leu Ala Ser Ala Asn Gly Asn Ser Val Val Val Lys Leu
421 422 423 424 425 426 427 428 429 430

Leu Leu Asp Asp Asp Lys Glu Leu Asn Val Leu Asp Asn Lys Lys Asp
431 432 433 434 435 436 437 438 439 440

His Ala Leu Ile Lys Ala Val Glu Lys Glu Glu Asp Glu Lys Ala Leu
441 442 443 444 445 446 447 448 449 450

Met Leu Leu Glu His Gly Thr Asp Leu Asn Ile Leu Asp Glu Tyr Gly
451 452 453 454 455 456 457 458 459 460

Asp Thr Thr Leu His Tyr Ala Ile Tyr Asn Glu Asp Lys Leu Met Ala
461 462 463 464 465 466 467 468 469 470

Lys Ala Leu Leu Leu Tyr Gly Ala Asp Ile Glu Ser Lys Asn Lys His
471 472 473 474 475 476 477 478 479 480

Gly Leu Thr Pro Leu Leu Leu Gly Val His Glu Glu Lys Glu Glu Val
481 482 483 484 485 486 487 488 489 490

Val Lys His Leu Ile Lys Lys Lys Ala Asp Leu Asn Ala Leu Asp Asp
491 492 493 494 495 496 497 498 499 500

Tyr Gly Asp Thr Ala Leu Ile Leu Ala Val Cys Cys Gly Ser Ala Ser
501 502 503 504 505 506 507 508 509 510

Ile Val Ser Leu Leu Leu Glu Glu Asp Ile Asp Val Ser Ser Glu Asp
511 512 513 514 515 516 517 518 519 520

Leu Ser Gly Glu Thr Ala Arg Glu Tyr Ala Val Ser Ser His His His
521 522 523 524 525 526 527 528 529 530

Val Ile Cys Glu Leu Leu Ser Asp Tyr Lys Glu Lys Glu Met Leu Lys
531 532 533 534 535 536 537 538 539 540

Ile Ser Ser Glu Asn Ser Asn Pro Glu Asn Val Ser Arg Thr Arg Asp
541 542 543 544 545 546 547 548 549 550

Lys

421 - 425

4211 - 4215

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